

## SEQUENCE LISTING

<110> Genentech, Inc. Ashkenazi, Avi Botstein, David Desnoyers, Luc Eaton, Dan L. Ferrara, Napoleone Filvaroff, Ellen Fong, Sherman Gao, Wei-Qiang Gerber, Hanspeter Gerritsen, Mary E. Goddard, A. Godowski, Paul J. Grimaldi, Christopher J. Gurney, Austin L. Hillan, Kenneth, J. Kljavin, Ivar J. Mather, Jennie P. Pan, James Paoni, Nicholas F. Roy, Margaret Ann Stewart, Timothy A. Tumas, Daniel Williams, P. Mickey Wood, William, I.

 $<\!\!120\!\!>$  Secreted and Transmembrane Polypeptides and Nucleic Acids Encoding the Same

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1825

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Ala Glu Tyr Phe Tyr Glu Phe Leu Ser Leu Arg Ser Leu Asp Lys Gly 100 105 110

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His Lys Ala Ser Val Val Gln Val Gly Phe Pro Cys Leu Gly Lys Gln 135 140

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Phe Tyr Pro Gly Lys Cys Ile Cys Pro Pro Gly Leu Glu Gly Glu Gln 260 265 270

Cys Glu Ile Ser Lys Cys Pro Gln Pro Cys Arg Asn Gly Gly Lys Cys 275 280 285

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Thr Asp Gln Leu Ser Arg Arg Gln Ile Arg Glu Tyr Gln Leu Tyr Ser 35 40 45

Arg Thr Ser Gly Lys His Val Gln Val Thr Gly Arg Arg Ile Ser Ala 50 55 60

Thr Ala Glu Asp Gly Asn Lys Phe Ala Lys Leu Ile Val Glu Thr Asp 65 70 75 80

Thr Phe Gly Ser Arg Val Arg Ile Lys Gly Ala Glu Ser Glu Lys Tyr 85 90 95

Ile Cys Met Asn Lys Arg Gly Lys Leu Ile Gly Lys Pro Ser Gly Lys
100 105 110

Ser Lys Asp Cys Val Phe Thr Glu Ile Val Leu Glu Asn Asn Tyr Thr 115 120 125

Ala Phe Gln Asn Ala Arg His Glu Gly Trp Phe Met Ala Phe Thr Arg 130 135 140

Gln Gly Arg Pro Arg Gln Ala Ser Arg Ser Arg Gln Asn Gln Arg Glu 145 150 155 160

Ala His Phe Ile Lys Arg Leu Tyr Gln Gly Gln Leu Pro Phe Pro Asn 165 170 175

His Ala Glu Lys Gln Lys Gln Phe Glu Phe Val Gly Ser Ala Pro Thr 180 185 190

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<213> Artificial Sequence

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<210> 27
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<212> DNA
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Cys Asn Glu Arg Ser Leu Thr Ser Val Pro Leu Gly Ile Pro Glu Gly 50

Val Thr Val Leu Tyr Leu His Asn Asn Gln Ile Asn Asn Ala Gly Phe 65

Pro Ala Glu Leu His Asn Val Gln Ser Val His Thr Val Tyr Leu Tyr 90

Gly Asn Gln Leu Asp Glu Phe Pro Met Asn Leu Pro Lys Asn Val Arg 110

Val Leu His Leu Gln Glu Asn Asn Ile Gln Thr Ile Ser Arg Ala Ala 115 120 125

Leu Ala Gln Leu Lys Leu Glu Glu Leu His Leu Asp Asp Asn Ser 130 135 Ile Ser Thr Val Gly Val Glu Asp Gly Ala Phe Arg Glu Ala Ile Ser 150 Leu Lys Leu Leu Phe Leu Ser Lys Asn His Leu Ser Ser Val Pro Val 170 Gly Leu Pro Val Asp Leu Gln Glu Leu Arg Val Asp Glu Asn Arg Ile 185 Ala Val Ile Ser Asp Met Ala Phe Gln Asn Leu Thr Ser Leu Glu Arg 200 Leu Ile Val Asp Gly Asn Leu Leu Thr Asn Lys Gly Ile Ala Glu Gly 215 Thr Phe Ser His Leu Thr Lys Leu Lys Glu Phe Ser Ile Val Arg Asn 230 235 Ser Leu Ser His Pro Pro Pro Asp Leu Pro Gly Thr His Leu Ile Arg 245 Leu Tyr Leu Gln Asp Asn Gln Ile Asn His Ile Pro Leu Thr Ala Phe 265 Ser Asn Leu Arg Lys Leu Glu Arg Leu Asp Ile Ser Asn Asn Gln Leu 280 Arg Met Leu Thr Gln Gly Val Phe Asp Asn Leu Ser Asn Leu Lys Gln 295 Leu Thr Ala Arg Asn Asn Pro Trp Phe Cys Asp Cys Ser Ile Lys Trp 310 315 Val Thr Glu Trp Leu Lys Tyr Ile Pro Ser Ser Leu Asn Val Arg Gly 325 Phe Met Cys Gln Gly Pro Glu Gln Val Arg Gly Met Ala Val Arg Glu 345 Leu Asn Met Asn Leu Leu Ser Cys Pro Thr Thr Pro Gly Leu Pro 355 360 Leu Phe Thr Pro Ala Pro Ser Thr Ala Ser Pro Thr Thr Gln Pro Pro 375 Thr Leu Ser Ile Pro Asn Pro Ser Arg Ser Tyr Thr Pro Pro Thr Pro 390 395 Thr Thr Ser Lys Leu Pro Thr Ile Pro Asp Trp Asp Gly Arg Glu Arg 405 Val Thr Pro Pro Ile Ser Glu Arg Ile Gln Leu Ser Ile His Phe Val 420 425

Asn Asp Thr Ser Ile Gln Val Ser Trp Leu Ser Leu Phe Thr Val Met 435 440 445

Ala Tyr Lys Leu Thr Trp Val Lys Met Gly His Ser Leu Val Gly Gly 450 . 455 . 460

Ile Val Gln Glu Arg Ile Val Ser Gly Glu Lys Gln His Leu Ser Leu 465 470 475 480

Val Asn Leu Glu Pro Arg Ser Thr Tyr Arg Ile Cys Leu Val Pro Leu
485 490 495

Asp Ala Phe Asn Tyr Arg Ala Val Glu Asp Thr Ile Cys Ser Glu Ala 500 505 510

Thr Thr His Ala Ser Tyr Leu Asn Asn Gly Ser Asn Thr Ala Ser Ser 515 520 525

His Glu Gln Thr Thr Ser His Ser Met Gly Ser Pro Phe Leu Leu Ala 530 535 540

Gly Leu Ile Gly Gly Ala Val Ile Phe Val Leu Val Val Leu Leu Ser 545 550 560

Val Phe Cys Trp His Met His Lys Lys Gly Arg Tyr Thr Ser Gln Lys 565 570 575

Trp Lys Tyr Asn Arg Gly Arg Arg Lys Asp Asp Tyr Cys Glu Ala Gly 580 585 590

Thr Lys Lys Asp Asn Ser Ile Leu Glu Met Thr Glu Thr Ser Phe Gln 595 600 605

Ile Val Ser Leu Asn Asn Asp Gln Leu Leu Lys Gly Asp Phe Arg Leu 610 615 620

Gln Pro Ile Tyr Thr Pro Asn Gly Gly Ile Asn Tyr Thr Asp Cys His 625 630 635 640

Ile Pro Asn Asn Met Arg Tyr Cys Asn Ser Ser Val Pro Asp Leu Glu 645 650 655

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<211> 21

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic oligonucleotide probe

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Ser Arg Gly Arg His Ala Arg Thr His Pro Gln Thr Ala Leu Leu Glu

20

35 40 45

Ser Ser Cys Glu Asn Lys Arg Ala Asp Leu Val Phe Ile Ile Asp Ser Ser Arg Ser Val Asn Thr His Asp Tyr Ala Lys Val Lys Glu Phe Ile Val Asp Ile Leu Gln Phe Leu Asp Ile Gly Pro Asp Val Thr Arg Val Gly Leu Leu Gln Tyr Gly Ser Thr Val Lys Asn Glu Phe Ser Leu Lys Thr Phe Lys Arg Lys Ser Glu Val Glu Arg Ala Val Lys Arg Met Arg 120 His Leu Ser Thr Gly Thr Met Thr Gly Leu Ala Ile Gln Tyr Ala Leu 130 135 Asn Ile Ala Phe Ser Glu Ala Glu Gly Ala Arg Pro Leu Arg Glu Asn Val Pro Arg Val Ile Met Ile Val Thr Asp Gly Arg Pro Gln Asp Ser 165 170 Val Ala Glu Val Ala Ala Lys Ala Arg Asp Thr Gly Ile Leu Ile Phe Ala Ile Gly Val Gly Gln Val Asp Phe Asn Thr Leu Lys Ser Ile Gly Ser Glu Pro His Glu Asp His Val Phe Leu Val Ala Asn Phe Ser Gln 210 215 Ile Glu Thr Leu Thr Ser Val Phe Gln Lys Lys Leu Cys Thr Ala His 230 Met Cys Ser Thr Leu Glu His Asn Cys Ala His Phe Cys Ile Asn Ile 245 Pro Gly Ser Tyr Val Cys Arg Cys Lys Gln Gly Tyr Ile Leu Asn Ser 265 Asp Gln Thr Thr Cys Arg Ile Gln Asp Leu Cys Ala Met Glu Asp His 275 280 Asn Cys Glu Gln Leu Cys Val Asn Val Pro Gly Ser Phe Val Cys Gln Cys Tyr Ser Gly Tyr Ala Leu Ala Glu Asp Gly Lys Arg Cys Val Ala 315 Val Asp Tyr Cys Ala Ser Glu Asn His Gly Cys Glu His Glu Cys Val 325 335 Asn Ala Asp Gly Ser Tyr Leu Cys Gln Cys His Glu Gly Phe Ala Leu

Asn Pro Asp Glu Lys Thr Cys Thr Arg Ile Asn Tyr Cys Ala Leu Asn 360 Lys Pro Gly Cys Glu His Glu Cys Val Asn Met Glu Glu Ser Tyr Tyr 375 Cys Arg Cys His Arg Gly Tyr Thr Leu Asp Pro Asn Gly Lys Thr Cys 395 Ser Arg Val Asp His Cys Ala Gln Gln Asp His Gly Cys Glu Gln Leu Cys Leu Asn Thr Glu Asp Ser Phe Val Cys Gln Cys Ser Glu Gly Phe 425 Leu Ile Asn Glu Asp Leu Lys Thr Cys Ser Arg Val Asp Tyr Cys Leu Leu Ser Asp His Gly Cys Glu Tyr Ser Cys Val Asn Met Asp Arg Ser 455 Phe Ala Cys Gln Cys Pro Glu Gly His Val Leu Arg Ser Asp Gly Lys 475 Thr Cys Ala Lys Leu Asp Ser Cys Ala Leu Gly Asp His Gly Cys Glu 485 His Ser Cys Val Ser Ser Glu Asp Ser Phe Val Cys Gln Cys Phe Glu 500 505 Gly Tyr Ile Leu Arg Glu Asp Gly Lys Thr Cys Arg Arg Lys Asp Val Cys Gln Ala Ile Asp His Gly Cys Glu His Ile Cys Val Asn Ser Asp 535 Asp Ser Tyr Thr Cys Glu Cys Leu Glu Gly Phe Arg Leu Ala Glu Asp 555 Gly Lys Arg Cys Arg Arg Lys Asp Val Cys Lys Ser Thr His His Gly 570 Cys Glu His Ile Cys Val Asn Asn Gly Asn Ser Tyr Ile Cys Lys Cys 580 Ser Glu Gly Phe Val Leu Ala Glu Asp Gly Arg Arg Cys Lys Lys Cys Thr Glu Gly Pro Ile Asp Leu Val Phe Val Ile Asp Gly Ser Lys Ser 615 Leu Gly Glu Glu Asn Phe Glu Val Val Lys Gln Phe Val Thr Gly Ile 625

Ile Asp Ser Leu Thr Ile Ser Pro Lys Ala Ala Arg Val Gly Leu Leu

645 650 655

Gln Tyr Ser Thr Gln Val His Thr Glu Phe Thr Leu Arg Asn Phe Asn 660 665 670

Ser Ala Lys Asp Met Lys Lys Ala Val Ala His Met Lys Tyr Met Gly 675 680 685

Lys Gly Ser Met Thr Gly Leu Ala Leu Lys His Met Phe Glu Arg Ser 690 695 700

Phe Thr Gln Gly Glu Gly Ala Arg Pro Leu Ser Thr Arg Val Pro Arg 705 710 715 720

Ala Ala Ile Val Phe Thr Asp Gly Arg Ala Gln Asp Asp Val Ser Glu
725 730 735

Trp Ala Ser Lys Ala Lys Ala Asn Gly Ile Thr Met Tyr Ala Val Gly 740 745 750

Val Gly Lys Ala Ile Glu Glu Glu Leu Gln Glu Ile Ala Ser Glu Pro
755 760 765

Thr Asn Lys His Leu Phe Tyr Ala Glu Asp Phe Ser Thr Met Asp Glu 770 780

Ile Ser Glu Lys Leu Lys Lys Gly Ile Cys Glu Ala Leu Glu Asp Ser
785 790 795 800

Asp Gly Arg Gln Asp Ser Pro Ala Gly Glu Leu Pro Lys Thr Val Gln 805 810 815

Gln Pro Thr Glu Ser Glu Pro Val Thr Ile Asn Ile Gln Asp Leu Leu 820 825 830

Ser Cys Ser Asn Phe Ala Val Gln His Arg Tyr Leu Phe Glu Glu Asp 835 840 845

Asn Leu Leu Arg Ser Thr Gln Lys Leu Ser His Ser Thr Lys Pro Ser 850 855 860

Gly Ser Pro Leu Glu Glu Lys His Asp Gln Cys Lys Cys Glu Asn Leu 865 870 875 880

Ile Met Phe Gln Asn Leu Ala Asn Glu Glu Val Arg Lys Leu Thr Gln 885 890 895

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Leu Ser Leu Thr Asn Leu Ser Ser Ser Met Ala Gly Val Tyr Val Cys 215 Lys Ala His Asn Glu Val Gly Thr Ala Gln Cys Asn Val Thr Leu Glu 230 235 225 Val Ser Thr Gly Pro Gly Ala Ala Val Val Ala Gly Ala Val Val Gly Thr Leu Val Gly Leu Gly Leu Leu Ala Gly Leu Val Leu Leu Tyr His 265 Arg Arg Gly Lys Ala Leu Glu Glu Pro Ala Asn Asp Ile Lys Glu Asp 280 Ala Ile Ala Pro Arg Thr Leu Pro Trp Pro Lys Ser Ser Asp Thr Ile 295 Ser Lys Asn Gly Thr Leu Ser Ser Val Thr Ser Ala Arg Ala Leu Arg 310 315 Pro Pro His Gly Pro Pro Arg Pro Gly Ala Leu Thr Pro Thr Pro Ser 330 325 Leu Ser Ser Gln Ala Leu Pro Ser Pro Arg Leu Pro Thr Thr Asp Gly 345 Ala His Pro Gln Pro Ile Ser Pro Ile Pro Gly Gly Val Ser Ser 360 Gly Leu Ser Arg Met Gly Ala Val Pro Val Met Val Pro Ala Gln Ser 375 Gln Ala Gly Ser Leu Val 385 <210> 40 <211> 22 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 40 agggtctcca ggagaaagac tc <210> 41 <211> 24 <212> DNA <213> Artificial Sequence

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Ser Tyr Tyr Cys Met Cys Val Pro Gly Phe Arg Ser Ser Ser Asn Gln 85 90 95

Asp Arg Phe Ile Thr Asn Asp Gly Thr Val Cys Ile Glu Asn Val Asn 100 105 110

Ala Asn Cys His Leu Asp Asn Val Cys Ile Ala Ala Asn Ile Asn Lys 115 120 125

Thr Leu Thr Lys Ile Arg Ser Ile Lys Glu Pro Val Ala Leu Leu Gln 130 135 140

Glu Val Tyr Arg Asn Ser Val Thr Asp Leu Ser Pro Thr Asp Ile Ile 145 150 155 160

Thr Tyr Ile Glu Ile Leu Ala Glu Ser Ser Ser Leu Leu Gly Tyr Lys 165 170 175

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180 185 190

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Leu His Lys Asn Phe Tyr Ile Phe Gly Tyr Leu Ser Pro Ala Val Val 530 535 540

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Val Cys Trp Leu Ser Thr Glu Asn Asn Phe Ile Trp Ser Phe Ile Gly 565 570 575

Pro Ala Cys Leu Ile Ile Leu Val Asn Leu Leu Ala Phe Gly Val Ile 580 585 590

Ile Tyr Lys Val Phe Arg His Thr Ala Gly Leu Lys Pro Glu Val Ser 595 600 ~ 605

Cys Phe Glu Asn Ile Arg Ser Cys Ala Arg Gly Ala Leu Ala Leu Leu 610 615 620

Phe Leu Leu Gly Thr Thr Trp Ile Phe Gly Val Leu His Val Val His 625 630 635 640

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Leu Ser Ser Ala Lys Gln Arg Gln Leu Tyr Lys Asn Arg Gly Phe Leu
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Lys Leu Gly Arg Ser Val Ser Phe Val Tyr Tyr Gln Gln Thr Leu Gln 65 70 75 80

Gly Asp Phe Lys Asn Arg Ala Glu Met Ile Asp Phe Asn Ile Arg Ile 85 90 95

Lys Asn Val Thr Arg Ser Asp Ala Gly Lys Tyr Arg Cys Glu Val Ser 100 105 110

Ala Pro Ser Glu Gln Gly Gln Asn Leu Glu Glu Asp Thr Val Thr Leu 115 120 125

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Ile Gly Leu His Asn Leu Leu Arg Leu His Leu Asn Ser Asn Arg Leu
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Lys Pro Leu Ile Asn Leu Arg Ser Leu Val Ile Ala Gly Ile Asn Leu
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Ile Ser Phe Tyr Asp Asn Arg Leu Ile Lys Val Pro His Val Ala Leu
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Thr Cys Ser Asn Ala Asn Leu Lys Glu Ile Pro Arg Asp Leu Pro Pro 50 55 60

Glu Thr Val Leu Leu Tyr Leu Asp Ser Asn Gln Ile Thr Ser Ile Pro 65 70 75 80

Asn Glu Ile Phe Lys Asp Leu His Gln Leu Arg Val Leu Asn Leu Ser 85 90 95

Lys Asn Gly Ile Glu Phe Ile Asp Glu His Ala Phe Lys Gly Val Ala 100 105 110

Glu Thr Leu Gln Thr Leu Asp Leu Ser Asp Asn Arg Ile Gln Ser Val 115 120 125

His Lys Asn Ala Phe Asn Asn Leu Lys Ala Arg Ala Arg Ile Ala Asn 130 135 140

Asn Pro Trp His Cys Asp Cys Thr Leu Gln Gln Val Leu Arg Ser Met
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Ala Ser Asn His Glu Thr Ala His Asn Val Ile Cys Lys Thr Ser Val 165 170 175

Leu Asp Glu His Ala Gly Arg Pro Phe Leu Asn Ala Ala Asn Asp Ala 180 185 190

Asp Leu Cys Asn Leu Pro Lys Lys Thr Thr Asp Tyr Ala Met Leu Val 195 200 205

Thr Met Phe Gly Trp Phe Thr Met Val Ile Ser Tyr Val Val Tyr Tyr 210 215 220

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Gln Val Phe Val Asp Glu Gly His Thr Val Gln Phe Val Cys Arg Ala 435 440 445

Asp Gly Asp Pro Pro Pro Ala Ile Leu Trp Leu Ser Pro Arg Lys His 450 455 460

Leu Val Ser Ala Lys Ser Asn Gly Arg Leu Thr Val Phe Pro Asp Gly 465 470 475 480

Thr Leu Glu Val Arg Tyr Ala Gln Val Gln Asp Asn Gly Thr Tyr Leu 485 490 495

Cys Ile Ala Ala Asn Ala Gly Gly Asn Asp Ser Met Pro Ala His Leu 500 505 510

His Val Arg Ser Tyr Ser Pro Asp Trp Pro His Gln Pro Asn Lys Thr 515 520 525

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Lys Cys Gln Val Lys Asp His Glu Asp Ser Ser Leu Gln Trp Ser Asn
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105

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120

115

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Thr Arg Ile Gln Glu Asp Pro Asn Gly Lys Thr Phe Thr Val Ser Ser 180 185 190

Ser Val Thr Phe Gln Val Thr Arg Glu Asp Asp Gly Ala Ser Ile Val 195 200 205

Cys Ser Val Asn His Glu Ser Leu Lys Gly Ala Asp Arg Ser Thr Ser 210 215 220

Gln Arg Ile Glu Val Leu Tyr Thr Pro Thr Ala Met Ile Arg Pro Asp 225 230 235 240

Pro Pro His Pro Arg Glu Gly Gln Lys Leu Leu His Cys Glu Gly
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Val Pro Pro Leu Lys Met Thr Gln Glu Ser Ala Leu Ile Phe Pro Phe 275 280 285

Leu Asn Lys Ser Asp Ser Gly Thr Tyr Gly Cys Thr Ala Thr Ser Asn 290 295 300

Met Gly Ser Tyr Lys Ala Tyr Tyr Thr Leu Asn Val Asn Asp Pro Ser 305 310 315 320

Pro Val Pro Ser Ser Ser Ser Thr Tyr His Ala Ile Ile Gly Gly Ile 325 330 335

Val Ala Phe Ile Val Phe Leu Leu Leu Ile Met Leu Ile Phe Leu Gly 340 345 350

His Tyr Leu Ile Arg His Lys Gly Thr Tyr Leu Thr His Glu Ala Lys 355 360 365

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45

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	agege eggtgtgage eagegetge				
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Phe Pro Gly Val Tyr Pro Pro Asn Ser Lys Cys Thr Trp Lys Ile Thr
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Val Pro Glu Gly Lys Val Val Leu Asn Phe Arg Phe Ile Asp Leu
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 65
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Glu Ser Asp Asn Leu Cys Arg Tyr Asp Phe Val Asp Val Tyr Asn Gly

His Ala Asn Gly Gln Arg Ile Gly Arg Phe Cys Gly Thr Phe Arg Pro

100

90

Asp Ala Asn Thr Ala Gly Asn Gly Phe Met Ala Met Phe Ser Ala Ala

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Asp Gly Arg Gly Lys Ile Met Pro Asn Ser Phe Ile Met Met Phe Lys 385 390 395

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375

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<sup>&</sup>lt;211> 22

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Pro Pro Pro Gln Ser Ser Pro Pro Pro Gln Pro His Pro Cys His Thr
Cys Arg Gly Leu Val Asp Ser Phe Asn Lys Gly Leu Glu Arg Thr Ile
Arg Asp Asn Phe Gly Gly Gly Asn Thr Ala Trp Glu Glu Glu Asn Leu
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                     70
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Ser Lys Tyr Lys Asp Ser Glu Thr Arg Leu Val Glu Val Leu Glu Gly
Val Cys Ser Lys Ser Asp Phe Glu Cys His Arg Leu Leu Glu Leu Ser
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Glu Glu Leu Val Glu Ser Trp Trp Phe His Lys Gln Gln Glu Ala Pro
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Asp Leu Phe Gln Trp Leu Cys Ser Asp Ser Leu Lys Leu Cys Cys Pro
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145
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Arg Pro Cys Gly Gly Tyr Gly Gln Cys Glu Gly Glu Gly Thr Arg Gly
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Gly Ser Gly His Cys Asp Cys Gln Ala Gly Tyr Gly Gly Glu Ala Cys
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Gly Gln Cys Gly Leu Gly Tyr Phe Glu Ala Glu Arg Asn Ala Ser His
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Glu Glu Ser Asn Cys Leu Gln Cys Lys Lys Gly Trp Ala Leu His His 225 230 235 240

Leu Lys Cys Val Asp Ile Asp Glu Cys Gly Thr Glu Gly Ala Asn Cys 245 250 255

Gly Ala Asp Gln Phe Cys Val Asn Thr Glu Gly Ser Tyr Glu Cys Arg 260 265 270

Asp Cys Ala Lys Ala Cys Leu Gly Cys Met Gly Ala Gly Pro Gly Arg 275 280 285

Cys Lys Lys Cys Ser Pro Gly Tyr Gln Gln Val Gly Ser Lys Cys Leu 290 295 300

Asp Val Asp Glu Cys Glu Thr Glu Val Cys Pro Gly Glu Asn Lys Gln 305 310 315 320

Cys Glu Asn Thr Glu Gly Gly Tyr Arg Cys Ile Cys Ala Glu Gly Tyr 325 330 335

Lys Gln Met Glu Gly Ile Cys Val Lys Glu Gln Ile Pro Glu Ser Ala 340 345 350

Gly Phe Phe Ser Glu Met Thr Glu Asp Glu Leu Val Val Leu Gln Gln 355 360 365

Met Phe Phe Gly Ile Ile Ile Cys Ala Leu Ala Thr Leu Ala Ala Lys 370 375 380

Gly Asp Leu Val Phe Thr Ala Ile Phe Ile Gly Ala Val Ala Ala Met 385 390 395 400

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<400> 114

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Glu Glu Leu Val Ile Pro Thr His Val Arg Ala Gln Tyr Val Ala Leu 50 55 60

Leu Gln Arg Ser His Gly Asp Arg Ser Arg Gly Lys Arg Phe Ser Gln 65 70 75 80

Ser Phe Arg Glu Val Ala Gly Arg Phe Leu Ala Leu Glu Ala Ser Thr 85 90 95

His Leu Leu Val Phe Gly Met Glu Gln Arg Leu Pro Pro Asn Ser Glu 100 105 110

Leu Val Gln Ala Val Leu Arg Leu Phe Gln Glu Pro Val Pro Lys Ala 115 120 125

Ala Leu His Arg His Gly Arg Leu Ser Pro Arg Ser Ala Arg Ala Arg 130 135 140

Val Thr Val Glu Trp Leu Arg Val Arg Asp Asp Gly Ser Asn Arg Thr 145 150 155 160

Ser Leu Ile Asp Ser Arg Leu Val Ser Val His Glu Ser Gly Trp Lys 165 170 175

Ala Phe Asp Val Thr Glu Ala Val Asn Phe Trp Gln Gln Leu Ser Arg 180 185 190

Pro Arg Gln Pro Leu Leu Gln Val Ser Val Gln Arg Glu His Leu 195 200 205

Gly Pro Leu Ala Ser Gly Ala His Lys Leu Val Arg Phe Ala Ser Gln 210 215 220

Gly Ala Pro Ala Gly Leu Gly Glu Pro Gln Leu Glu Leu His Thr Leu 225 230 235 240

Asp Leu Gly Asp Tyr Gly Ala Gln Gly Asp Cys Asp Pro Glu Ala Pro 245 250 255

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Gln Gly Met Lys Trp Ala Glu Asn Trp Val Leu Glu Pro Pro Gly Phe 275 280 285				
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Ala Phe Lys Trp Pro Phe Leu Gly Pro Arg Gln Cys Ile Ala Ser Glu 305 310 315 320				
Thr Asp Ser Leu Pro Met Ile Val Ser Ile Lys Glu Gly Gly Arg Thr 325 330 335				
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                                                 45
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Ser Cys Ala Tyr Ser Gly Phe Ser Ser Pro Arg Val Glu Trp Lys Phe
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Lys Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met Val Ser 100 105 110

Glu Glu Gly Gly Asn Ser Tyr Gly Glu Val Lys Val Lys Leu Ile Val 115 120 125

Leu Val Pro Pro Ser Lys Pro Thr Val Asn Ile Pro Ser Ser Ala Thr 130 135 140

Ile Gly Asn Arg Ala Val Leu Thr Cys Ser Glu Gln Asp Gly Ser Pro 145 150 155 160

Pro Ser Glu Tyr Thr Trp Phe Lys Asp Gly Ile Val Met Pro Thr Asn 165 170 175

Pro Lys Ser Thr Arg Ala Phe Ser Asn Ser Ser Tyr Val Leu Asn Pro 180 185 190

Thr Thr Gly Glu Leu Val Phe Asp Pro Leu Ser Ala Ser Asp Thr Gly 195 200 205

Glu Tyr Ser Cys Glu Ala Arg Asn Gly Tyr Gly Thr Pro Met Thr Ser 210 215 220

Asn Ala Val Arg Met Glu Ala Val Glu Arg Asn Val Gly Val Ile Val 225 230 235 240

Ala Ala Val Leu Val Thr Leu Ile Leu Cly Ile Leu Val Phe Cly 245 250 255

Ile Trp Phe Ala Tyr Ser Arg Gly His Phe Asp Arg Thr Lys Lys Gly 260 265 270

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        35
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                                                 45
Pro Ser Ser Gly Ser Cys Pro Pro Thr Lys Phe Gln Cys Arg Thr Ser
Gly Leu Cys Val Pro Leu Thr Trp Arg Cys Asp Arg Asp Leu Asp Cys
65
                     70
                                         75
Ser Asp Gly Ser Asp Glu Glu Glu Cys Arg Ile Glu Pro Cys Thr Gln
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Lys Gly Gln Cys Pro Pro Pro Pro Gly Leu Pro Cys Pro Cys Thr Gly

100 105 110

Val Ser Asp Cys Ser Gly Gly Thr Asp Lys Leu Arg Asn Cys Ser 115 120 125

Arg Leu Ala Cys Leu Ala Gly Glu Leu Arg Cys Thr Leu Ser Asp Asp 130 135 140

Cys Ile Pro Leu Thr Trp Arg Cys Asp Gly His Pro Asp Cys Pro Asp 145 150 155 160

Ser Ser Asp Glu Leu Gly Cys Gly Thr Asn Glu Ile Leu Pro Glu Gly 165 170 175

Asp Ala Thr Thr Met Gly Pro Pro Val Thr Leu Glu Ser Val Thr Ser 180 185 190

Leu Arg Asn Ala Thr Thr Met Gly Pro Pro Val Thr Leu Glu Ser Val 195 200 205

Pro Ser Val Gly Asn Ala Thr Ser Ser Ser Ala Gly Asp Gln Ser Gly 210 215 220

Ser Pro Thr Ala Tyr Gly Val Ile Ala Ala Ala Ala Val Leu Ser Ala 225 230 235 240

Ser Leu Val Thr Ala Thr Leu Leu Leu Leu Ser Trp Leu Arg Ala Gln 245 250 255

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<213> Artificial Sequence

<220>

<400> 128

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24

<210> 129

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<400> 129

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<210> 131
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265

Glu Phe Tyr Cys Asp Pro Gly Tyr Ser Leu Thr Ser Asp Tyr Lys Tyr 275 280 285

Ile Thr Cys Gln Tyr Gly Glu Trp Phe Pro Ser Tyr Gln Val Tyr Cys 290 295 300

Ile Lys Ser Glu Gln Thr Trp Pro Ser Thr His Glu Thr Leu Leu Thr 305 310 315 320

Thr Trp Lys Ile Val Ala Phe Thr Ala Thr Ser Val Leu Leu Val Leu 325 330 335

Leu Leu Val Ile Leu Ala Arg Met Phe Gln Thr Lys Phe Lys Ala His  $340 \hspace{1.5cm} 345 \hspace{1.5cm} 350$ 

Phe Pro Pro Arg Gly Pro Pro Arg Ser Ser Ser Ser Asp Pro Asp Phe 355 360 365

Val Val Val Asp Gly Val Pro Val Met Leu Pro Ser Tyr Asp Glu Ala 370 375 380

Val Ser Gly Gly Leu Ser Ala Leu Gly Pro Gly Tyr Met Ala Ser Val 385 390 395 400

Gly Gln Gly Cys Pro Leu Pro Val Asp Asp Gln Ser Pro Pro Ala Tyr 405 410 415

Pro Gly Ser Gly Asp Thr Asp Thr Gly Pro Gly Glu Ser Glu Thr Cys 420 425 430

Asp Ser Val Ser Gly Ser Ser Glu Leu Leu Gln Ser Leu Tyr Ser Pro 435 440 445

Pro Arg Cys Gln Glu Ser Thr His Pro Ala Ser Asp Asn Pro Asp Ile 450 455 460

Ile Ala Ser Thr Ala Glu Glu Val Ala Ser Thr Ser Pro Gly Ile His 465 470 475 480

His Ala His Trp Val Leu Phe Leu Arg Asn 485 490

<210> 133

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
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<400> 133

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<210> 134

<211> 23

<212> DNA

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<212> DNA
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<212> DNA
<213> Homo sapiens
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<212> PRT

<213> Homo sapiens

<400> 137

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20 25 30

Asp Leu Arg Gly Gly Gln Pro Val Cys Arg Gly Gly Thr Gln Arg Pro 35 40 45

Cys Tyr Lys Val Ile Tyr Phe His Asp Thr Ser Arg Arg Leu Asn Phe 50 55 60

Glu Glu Ala Lys Glu Ala Cys Arg Arg Asp Gly Gly Gln Leu Val Ser 65 70 75 80

Ile Glu Ser Glu Asp Glu Gln Lys Leu Ile Glu Lys Phe Ile Glu Asn 85 90 95

Leu Leu Pro Ser Asp Gly Asp Phe Trp Ile Gly Leu Arg Arg Glu
100 105 110

Glu Lys Gln Ser Asn Ser Thr Ala Cys Gln Asp Leu Tyr Ala Trp Thr 115 120 125

Asp Gly Ser Ile Ser Gln Phe Arg Asn Trp Tyr Val Asp Glu Pro Ser 130 135 140

Cys Gly Ser Glu Val Cys Val Val Met Tyr His Gln Pro Ser Ala Pro 145 150 155 160

Ala Gly Ile Gly Gly Pro Tyr Met Phe Gln Trp Asn Asp Asp Arg Cys 165 170 175

Asn Met Lys Asn Asn Phe Ile Cys Lys Tyr Ser Asp Glu Lys Pro Ala 180 185 190

Val Pro Ser Arg Glu Ala Glu Gly Glu Glu Thr Glu Leu Thr Thr Pro 195 200 205

Val Leu Pro Glu Glu Thr Gln Glu Glu Asp Ala Lys Lys Thr Phe Lys 210 215 220

Glu Ser Arg Glu Ala Ala Leu Asn Leu Ala Tyr Ile Leu Ile Pro Ser 225 230 235 240

Ile Pro Leu Leu Leu Leu Val Val Thr Thr Val Val Cys Trp Val 245 250 255

Trp Ile Cys Arg Lys Arg Lys Arg Glu Gln Pro Asp Pro Ser Thr Lys

260 265 270

Lys Gln His Thr Ile Trp Pro Ser Pro His Gln Gly Asn Ser Pro Asp 275 280 285

Leu Glu Val Tyr Asn Val Ile Arg Lys Gln Ser Glu Ala Asp Leu Ala 290 295 300

Glu Thr Arg Pro Asp Leu Lys Asn Ile Ser Phe Arg Val Cys Ser Gly 315 310 315 320

Glu Ala Thr Pro Asp Asp Met Ser Cys Asp Tyr Asp Asn Met Ala Val\$325\$ 330 335

Asn Pro Ser Glu Ser Gly Phe Val Thr Leu Val Ser Val Glu Ser Gly 340 345 350

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Ser Lys Glu Ser Gly Trp Val Glu Asn Glu Ile Tyr Gly Tyr 370 375 380

<210> 138

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

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<210> 139

<211> 24

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<213> Artificial Sequence

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<400> 139

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24

<210> 140

<211> 24

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
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<211> 1514
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<213> Homo sapiens
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<211> 428
<212> PRT
<213> Homo sapiens
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             20
                                 25
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Lys Leu Lys Met Val Gln Val Val Phe Arg His Gly Ala Arg Ser Pro
                         55
Leu Lys Pro Leu Pro Leu Glu Glu Gln Val Glu Trp Asn Pro Gln Leu
 65
Leu Glu Val Pro Pro Gln Thr Gln Phe Asp Tyr Thr Val Thr Asn Leu
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90 95

Ala Gly Gly Pro Lys Pro Tyr Ser Pro Tyr Asp Ser Gln Tyr His Glu Thr Thr Leu Lys Gly Gly Met Phe Ala Gly Gln Leu Thr Lys Val Gly 120 Met Gln Gln Met Phe Ala Leu Gly Glu Arg Leu Arg Lys Asn Tyr Val 135 Glu Asp Ile Pro Phe Leu Ser Pro Thr Phe Asn Pro Gln Glu Val Phe 150 155 Ile Arg Ser Thr Asn Ile Phe Arg Asn Leu Glu Ser Thr Arg Cys Leu 165 Leu Ala Gly Leu Phe Gln Cys Gln Lys Glu Gly Pro Ile Ile His 185 Thr Asp Glu Ala Asp Ser Glu Val Leu Tyr Pro Asn Tyr Gln Ser Cys Trp Ser Leu Arg Gln Arg Thr Arg Gly Arg Arg Gln Thr Ala Ser Leu 215 Gln Pro Gly Ile Ser Glu Asp Leu Lys Lys Val Lys Asp Arg Met Gly 235 230 Ile Asp Ser Ser Asp Lys Val Asp Phe Phe Ile Leu Leu Asp Asn Val 245 250 Ala Ala Glu Gln Ala His Asn Leu Pro Ser Cys Pro Met Leu Lys Arg 265 Phe Ala Arg Met Ile Glu Gln Arg Ala Val Asp Thr Ser Leu Tyr Ile 275 Leu Pro Lys Glu Asp Arg Glu Ser Leu Gln Met Ala Val Gly Pro Phe 295 Leu His Ile Leu Glu Ser Asn Leu Leu Lys Ala Met Asp Ser Ala Thr 315 Ala Pro Asp Lys Ile Arg Lys Leu Tyr Leu Tyr Ala Ala His Asp Val Thr Phe Ile Pro Leu Leu Met Thr Leu Gly Ile Phe Asp His Lys Trp 345 Pro Pro Phe Ala Val Asp Leu Thr Met Glu Leu Tyr Gln His Leu Glu Ser Lys Glu Trp Phe Val Gln Leu Tyr Tyr His Gly Lys Glu Gln Val 375 Pro Arg Gly Cys Pro Asp Gly Leu Cys Pro Leu Asp Met Phe Leu Asn

Ala Met Ser Val Tyr Thr Leu Ser Pro Glu Lys Tyr His Ala Leu Cys 405  Ser Gln Thr Gln Val Met Glu Val Gly Asn Glu Glu 420  425  <210> 143 <211> 24 <212> DNA <213> Artificial Sequence  <220> <223> Description of Artificial Sequence: Synthetic
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2223 Description of Artificial Sequence: Synthetic
oligonucleotide probe
<400> 143
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<223> Description of Artificial Sequence: Synthetic oligonucleotide probe
<400> 144
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<210> 145
<211> 24
<212> DNA
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<223> Description of Artificial Sequence: Synthetic oligonucleotide probe
<400> 145
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<210> 146
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<220>
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ggcagagaac cagaggccgg aggagactgc ctctttacag ccagg
<210> 147

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90 95

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Ser	Ser 130	Phe	Ser	Pro	Val	Pro 135	Glu	Gly	Val	Arg	Leu 140	Ala	Asp	Gly	Pro
Gly 145	His	Cys	Lys	Gly	Arg 150	Val	Glu	Val	Lys	His 155	Gln	Asn	Gln	Trp	Tyr 160
Thr	Val	Cys	Gln	Thr 165	Gly	Trp	Ser	Leu	Arg 170	Ala	Ala	Lys	Val	Val 175	Cys
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Lys	His	Ala 195	Tyr	Gly	Arg	Lys	Pro 200	Ile	Trp	Leu	Ser	Gln 205	Met	Ser	Cys
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Leu	Glu	Val	Leu 260	His	Lys	Gly	Val	Trp 265	Gly	Ser	Val	Суѕ	Asp 270	Asp	Asn
Trp	Gly	Glu 275	Lys	Glu	Asp	Gln	Val 280	Val	Cys	Lys	Gln	Leu 285	Gly	Cys	Gly
Lys	Ser 290	Leu	Ser	Pro	Ser	Phe 295	Arg	Asp	Arg	Lys	Cys 300	Tyr	Gly	Pro	Gly
Val 305	Gly	Arg	Ile	Trp	Leu 310	Asp	Asn	Val	Arg	Cys 315	Ser	Gly	Glu	Glu	Gln 320
Ser	Leu	Glu	Gln	Cys 325	Gln	His	Arg	Phe	Trp 330	Gly	Phe	His	Asp	Cys 335	Thr
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Ala Tyr Leu Arg Asn Ala Val Val Val Ile Thr Gly Ala Thr Ser Gly 35 40 45

Leu Gly Lys Glu Cys Ala Lys Val Phe Tyr Ala Ala Gly Ala Lys Leu 50 55 60

Val Leu Cys Gly Arg Asn Gly Gly Ala Leu Glu Glu Leu Ile Arg Glu
65 70 75 80

Leu Thr Ala Ser His Ala Thr Lys Val Gln Thr His Lys Pro Tyr Leu 85 90 95

Val Thr Phe Asp Leu Thr Asp Ser Gly Ala Ile Val Ala Ala Ala 100 105 110

Glu Ile Leu Gln Cys Phe Gly Tyr Val Asp Ile Leu Val Asn Asn Ala 115 120 125

Gly Ile Ser Tyr Arg Gly Thr Ile Met Asp Thr Thr Val Asp Val Asp 130 135 140

Lys Arg Val Met Glu Thr Asn Tyr Phe Gly Pro Val Ala Leu Thr Lys 145 150 155 160

Ala Leu Leu Pro Ser Met Ile Lys Arg Arg Gln Gly His Ile Val Ala 165 170 175

Ile Ser Ser Ile Gln Gly Lys Met Ser Ile Pro Phe Arg Ser Ala Tyr 180 185 190

Ala Ala Ser Lys His Ala Thr Gln Ala Phe Phe Asp Cys Leu Arg Ala 195 200 205

Glu Met Glu Gln Tyr Glu Ile Glu Val Thr Val Ile Ser Pro Gly Tyr 210 215 220

Ile His Thr Asn Leu Ser Val Asn Ala Ile Thr Ala Asp Gly Ser Arg 225 230 235 240

Tyr Gly Val Met Asp Thr Thr Thr Ala Gln Gly Arg Ser Pro Val Glu 245 250 255

Val Ala Gln Asp Val Leu Ala Ala Val Gly Lys Lys Lys Asp Val Ile Leu Ala Asp Leu Leu Pro Ser Leu Ala Val Tyr Leu Arg Thr Leu Ala Pro Gly Leu Phe Phe Ser Leu Met Ala Ser Arg Ala Arg Lys Glu 295 290 300 Arg Lys Ser Lys Asn Ser 305 <210> 154 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 154 24 ggtgctaaac tggtgctctg tggc <210> 155 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 155 cagggcaaga tgagcattcc 20 <210> 156 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 156 tcatactgtt ccatctcggc acgc 24 <210> 157 <211> 50 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe

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Ser Val Thr Gly Glu Ile Val Leu Ile Thr Gly Ala Gly His Gly Ile
Gly Arg Leu Thr Ala Tyr Glu Phe Ala Lys Leu Lys Ser Lys Leu Val
                       55
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Leu Trp Asp Ile Asn Lys His Gly Leu Glu Glu Thr Ala Ala Lys Cys
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Lys Gly Leu Gly Ala Lys Val His Thr Phe Val Val Asp Cys Ser Asn 85 90 95

Arg Glu Asp Ile Tyr Ser Ser Ala Lys Lys Val Lys Ala Glu Ile Gly 100 105 110

Asp Val Ser Ile Leu Val Asn Asn Ala Gly Val Val Tyr Thr Ser Asp 115 120 125

Leu Phe Ala Thr Gln Asp Pro Gln Ile Glu Lys Thr Phe Glu Val Asn 130 135 140

Val Leu Ala His Phe Trp Thr Thr Lys Ala Phe Leu Pro Ala Met Thr 145 150 155 160

Lys Asn Asn His Gly His Ile Val Thr Val Ala Ser Ala Ala Gly His 165 170 175

Val Ser Val Pro Phe Leu Leu Ala Tyr Cys Ser Ser Lys Phe Ala Ala 180 185 190

Val Gly Phe His Lys Thr Leu Thr Asp Glu Leu Ala Ala Leu Gln Ile 195 200 205

Thr Gly Val Lys Thr Thr Cys Leu Cys Pro Asn Phe Val Asn Thr Gly 210 215 220

Phe Ile Lys Asn Pro Ser Thr Ser Leu Gly Pro Thr Leu Glu Pro Glu 225 230 235 240

Glu Val Val Asn Arg Leu Met His Gly Ile Leu Thr Glu Gln Lys Met 245 250 255

Ile Phe Ile Pro Ser Ser Ile Ala Phe Leu Thr Thr Leu Glu Arg Ile 260 265 270

Leu Pro Glu Arg Phe Leu Ala Val Leu Lys Arg Lys Ile Ser Val Lys 275 280 285

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<223> Description of Artificial Sequence: Synthetic
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Thr 145	Leu	Ser	Met	Leu	Tyr 150	Ile	Asp	Asn	Pro	Val 155	Gly	Thr	Gly	Phe	Ser 160	
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Arg	Asp	Leu	Tyr 180	Ser	Ala	Leu	Ile	Gln 185	Phe	Phe	Gln	Ile	Phe 190	Pro	Glu	
Tyr	Lys	Asn 195	Asn	Asp	Phe	Tyr	Val 200	Thr	Gly	Glu	Ser	Tyr 205	Ala	Gly	Lys	
Tyr	Val	Pro	Ala	Ile	Ala	His	Leu	Ile	His	Ser	Leu	Asn	Pro	Val	Arg	

Glu Val Lys Ile Asn Leu Asn Gly Ile Ala Ile Gly Asp Gly Tyr Ser 225 230 235 240

Asp Pro Glu Ser Ile Ile Gly Gly Tyr Ala Glu Phe Leu Tyr Gln Ile
245 250 . 255

Gly Leu Leu Asp Glu Lys Gln Lys Lys Tyr Phe Gln Lys Gln Cys His 260 265 270

Glu Cys Ile Glu His Ile Arg Lys Gln Asn Trp Phe Glu Ala Phe Glu 275 280 285

Ile Leu Asp Lys Leu Leu Asp Gly Asp Leu Thr Ser Asp Pro Ser Tyr 290 295 300

Phe Gln Asn Val Thr Gly Cys Ser Asn Tyr Tyr Asn Phe Leu Arg Cys 305 310 315 320

Thr Glu Pro Glu Asp Gln Leu Tyr Tyr Val Lys Phe Leu Ser Leu Pro 325 330 335

Glu Val Arg Gln Ala Ile His Val Gly Asn Gln Thr Phe Asn Asp Gly 340 345 350

Thr Ile Val Glu Lys Tyr Leu Arg Glu Asp Thr Val Gln Ser Val Lys 355 360 365

Pro Trp Leu Thr Glu Ile Met Asn Asn Tyr Lys Val Leu Ile Tyr Asn 370 375 380

Gly Gln Leu Asp Ile Ile Val Ala Ala Ala Leu Thr Glu Arg Ser Leu 385 390 395 400

Met Gly Met Asp Trp Lys Gly Ser Gln Glu Tyr Lys Lys Ala Glu Lys 405 410 415

Lys Val Trp Lys Ile Phe Lys Ser Asp Ser Glu Val Ala Gly Tyr Ile 420 425 430

Arg Gln Ala Gly Asp Phe His Gln Val Ile Ile Arg Gly Gly His 435 440 445

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<212> DNA

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<223> Description of Artificial Sequence: Synthetic
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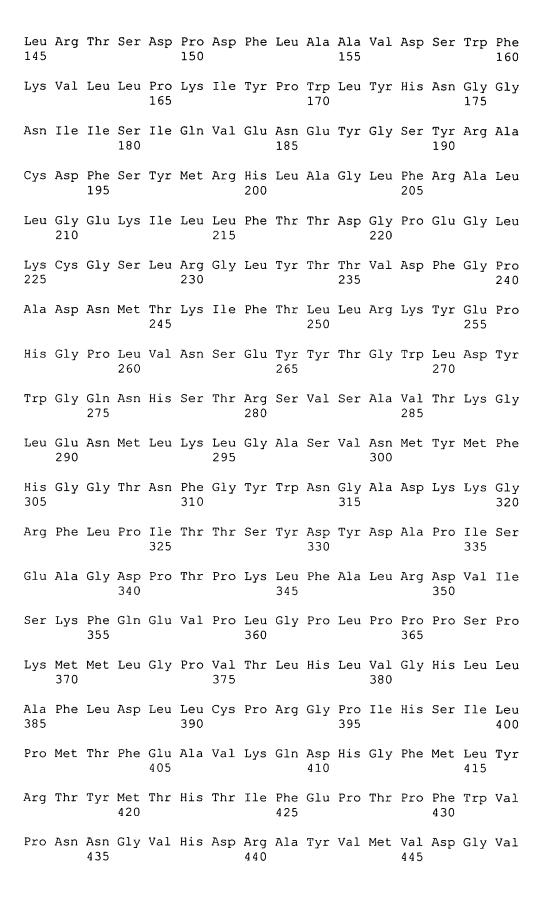
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Glu Met Gly Gly Leu Pro Ser Trp Leu Leu Arg Lys Pro Glu Ile His 135 130 140



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370 375 380

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His Ile Thr Gly Asn Lys Val Asp Ile Leu Pro Lys Gln Leu Phe Lys
405 410 415

Cys Ile Lys Leu Arg Thr Leu Asn Leu Gly Gln Asn Cys Ile Thr Ser 420 425 430

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Lys Gly Asn Cys Leu Asp Arg Leu Pro Ala Gln Leu Gly Gln Cys Arg 450 455 460

Met Leu Lys Lys Ser Gly Leu Val Val Glu Asp His Leu Phe Asp Thr 465 470 475 480

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Pro Glu Asn Lys Ser Ile Arg Ile Ile Phe Ser Tyr Val Gln Leu Asp 65 70 75 80

Pro Asp Gly Ser Cys Glu Ser Glu Asn Ile Lys Val Phe Asp Gly Thr 85 90 95

Ser Ser Asn Gly Pro Leu Leu Gly Gln Val Cys Ser Lys Asn Asp Tyr 100 105 110

Val Pro Val Phe Glu Ser Ser Ser Ser Thr Leu Thr Phe Gln Ile Val 115 120 125

Thr Asp Ser Ala Arg Ile Gln Arg Thr Val Phe Val Phe Tyr Tyr Phe 130 135 140

Phe Ser Pro Asn Ile Ser Ile Pro Asn Cys Gly Gly Tyr Leu Asp Thr 145 150 155 160

Leu Glu Gly Ser Phe Thr Ser Pro Asn Tyr Pro Lys Pro His Pro Glu 165 170 175

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Lys Leu Asn Phe Lys Glu Ile Phe Leu Glu Ile Asp Lys Gln Cys Lys 195 200 205

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Asn Ser Leu Thr Val Val Leu Ser Thr Asp Tyr Ala Asn Ser Tyr Arg 245 250 255

Gly Phe Ser Ala Ser Tyr Thr Ser Ile Tyr Ala Glu Asn Ile Asn Thr 260 265 Thr Ser Leu Thr Cys Ser Ser Asp Arg Met Arg Val Ile Ile Ser Lys Ser Tyr Leu Glu Ala Phe Asn Ser Asn Gly Asn Asn Leu Gln Leu Lys 295 Asp Pro Thr Cys Arg Pro Lys Leu Ser Asn Val Val Glu Phe Ser Val 310 315 Pro Leu Asn Gly Cys Gly Thr Ile Arg Lys Val Glu Asp Gln Ser Ile 330 Thr Tyr Thr Asn Ile Ile Thr Phe Ser Ala Ser Ser Thr Ser Glu Val Ile Thr Arg Gln Lys Gln Leu Gln Ile Ile Val Lys Cys Glu Met Gly 360 His Asn Ser Thr Val Glu Ile Ile Tyr Ile Thr Glu Asp Asp Val Ile 370 Gln Ser Gln Asn Ala Leu Gly Lys Tyr Asn Thr Ser Met Ala Leu Phe 390 Glu Ser Asn Ser Phe Glu Lys Thr Ile Leu Glu Ser Pro Tyr Tyr Val 405 410 Asp Leu Asn Gln Thr Leu Phe Val Gln Val Ser Leu His Thr Ser Asp 420 Pro Asn Leu Val Val Phe Leu Asp Thr Cys Arg Ala Ser Pro Thr Ser 440 Asp Phe Ala Ser Pro Thr Tyr Asp Leu Ile Lys Ser Gly Cys Ser Arg 455 Asp Glu Thr Cys Lys Val Tyr Pro Leu Phe Gly His Tyr Gly Arg Phe 465 470 Gln Phe Asn Ala Phe Lys Phe Leu Arg Ser Met Ser Ser Val Tyr Leu 490 Gln Cys Lys Val Leu Ile Cys Asp Ser Ser Asp His Gln Ser Arg Cys Asn Gln Gly Cys Val Ser Arg Ser Lys Arg Asp Ile Ser Ser Tyr Lys 520 Trp Lys Thr Asp Ser Ile Ile Gly Pro Ile Arg Leu Lys Arg Asp Arg 530 Ser Ala Ser Gly Asn Ser Gly Phe Gln His Glu Thr His Ala Glu Glu 545 550 555

Thr Pro Asn Gln Pro Phe Asn Ser Val His Leu Phe Ser Phe Met Val 565 570 575 Leu Ala Leu Asn Val Val Thr Val Ala Thr Ile Thr Val Arg His Phe 580 585 590 Val Asn Gln Arg Ala Asp Tyr Lys Tyr Gln Lys Leu Gln Asn Tyr 600 <210> 191 <211> 21 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 191 tctctattcc aaactgtggc g 21 <210> 192 <211> 22 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic oligonucleotide probe <400> 192 tttgatgacg attcgaaggt gg 22 <210> 193 <211> 47 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic oligonucleotide probe ggaaggatcc ttcaccagcc ccaattaccc aaagccgcat cctgagc 47 <210> 194 <211> 2362 <212> DNA <213> Homo sapiens <400> 194 gacggaagaa cagcgctccc gaggccgcgg gagcctgcag agaggacagc cggcctgcgc 60 cgggacatgc ggccccagga gctccccagg ctcgcgttcc cgttgctgct gttgctgttg 120 ctgctgctgc cgccgccgcc gtgccctgcc cacagcgcca cgcgcttcga ccccacctgg 180 gagteectgg aegeeegeea getgeeegeg tggtttgaee aggeeaagtt eggeatette 240 atccactggg gagtgttttc cgtgcccagc ttcggtagcg agtggttctg gtggtattgg 300

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Gly Leu Ser Leu Ala Gly Lys Asp Gln Val Glu Thr Ala Leu Lys Ala 65 70 75 80

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Glu Phe Ile Val Ser Asp Ser Thr Tyr Ser Val Ala Ser Pro Tyr Ser 165 170 175

Thr Ile Pro Ala Pro Thr Thr Pro Pro Ala Pro Ala Ser Thr Ser 180 185 190

Ile Pro Arg Arg Lys Lys Leu Ile Cys Val Thr Glu Val Phe Met Glu 195 200 205

Thr Ser Thr Met Ser Thr Glu Thr Glu Pro Phe Val Glu Asn Lys Ala 210 215 220

Ala Phe Lys Asn Glu Ala Ala Gly Phe Gly Gly Val Pro Thr Ala Leu 225 230 235 240

Leu Val Leu Ala Leu Leu Phe Phe Gly Ala Ala Ala Gly Leu Gly Phe 245 250 255

Cys Tyr Val Lys Arg Tyr Val Lys Ala Phe Pro Phe Thr Asn Lys Asn

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Leu Pro Thr Gln Arg Glu Asp Gly Asn Pro Cys Asp Phe Asp Trp Arg 50 55 60

Glu Val Glu Ile Leu Met Phe Leu Ser Ala Ile Val Met Met Lys Asn 65 70 75 80

Arg Arg Ser Ile Thr Val Glu Gln His Ile Gly Asn Ile Phe Met Phe  $85 \hspace{1.5cm} 90 \hspace{1.5cm} 95$ 

Ser Lys Val Ala Asn Thr Ile Leu Phe Phe Arg Leu Asp Ile Arg Met 100 105 110

Gly Leu Leu Tyr Ile Thr Leu Cys Ile Val Phe Leu Met Thr Cys Lys 115 120 125

Pro Pro Leu Tyr Met Gly Pro Glu Tyr Ile Lys Tyr Phe Asn Asp Lys 130 135 140

Thr Ile Asp Glu Glu Leu Glu Arg Asp Lys Arg Val Thr Trp Ile Val 145 150 155 160

Glu Phe Phe Ala Asn Trp Ser Asn Asp Cys Gln Ser Phe Ala Pro Ile 165 170 175

Tyr Ala Asp Leu Ser Leu Lys Tyr Asn Cys Thr Gly Leu Asn Phe Gly
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Lys Val Asp Val Gly Arg Tyr Thr Asp Val Ser Thr Arg Tyr Lys Val 195 200 205

Ser Thr Ser Pro Leu Thr Lys Gln Leu Pro Thr Leu Ile Leu Phe Gln 210 215 220

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Val	Ser 130	Leu	Glu	Ile	Gln	Asp 135	Leu	Arg	Leu	Glu	Asp 140	Tyr	Gly	Arg	Tyr
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Leu Arg Ser Ala Val Glu Glu Met Glu Ala Glu Glu Ala Ala Lys
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Ala Ser Ser Glu Val Asn Leu Ala Asn Leu Pro Pro Ser Tyr His Asn 85 90 95

Glu Thr Asn Thr Asp Thr Lys Val Gly Asn Asn Thr Ile His Val His 100 105 110

Arg Glu Ile His Lys Ile Thr Asn Asn Gln Thr Gly Gln Met Val Phe 115 120 125

Ser Glu Thr Val Ile Thr Ser Val Gly Asp Glu Glu Gly Arg Arg Ser 130 135 140

His Glu Cys Ile Ile Asp Glu Asp Cys Gly Pro Ser Met Tyr Cys Gln 145 150 155 160

Phe Ala Ser Phe Gln Tyr Thr Cys Gln Pro Cys Arg Gly Gln Arg Met 165 170 175

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Gly His Cys Thr Lys Met Ala Thr Arg Gly Ser Asn Gly Thr Ile Cys 195 200 205

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Gly Leu Leu Phe Pro Val Cys Thr Pro Leu Pro Val Glu Gly Glu Leu 225 230 235 240

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Ser Tyr Arg Glu Ala Thr Thr Val Asp Cys Asn Asp Leu Phe Leu
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Ser Asp Ala Arg Asp Cys Asp Phe His Ala Leu Pro Gln Leu Leu 110 115 120

Ser Leu His Leu Glu Glu Asn Gln Leu Thr Arg Leu Glu Asp His 125 130 135

Ser Phe Ala Gly Leu Ala Ser Leu Gln Glu Leu Tyr Leu Asn His 140 145 150 Asn Gln Leu Tyr Arg Ile Ala Pro Arg Ala Phe Ser Gly Leu Ser Asn Leu Leu Arg Leu His Leu Asn Ser Asn Leu Leu Arg Ala Ile Asp Ser Arg Trp Phe Glu Met Leu Pro Asn Leu Glu Ile Leu Met Ile Gly Gly Asn Lys Val Asp Ala Ile Leu Asp Met Asn Phe Arg Pro Leu Ala Asn Leu Arg Ser Leu Val Leu Ala Gly Met Asn Leu Arg Glu Ile Ser Asp Tyr Ala Leu Glu Gly Leu Gln Ser Leu Glu Ser Leu Ser Phe Tyr Asp Asn Gln Leu Ala Arg Val Pro Arg Arg Ala Leu Glu Gln Val Pro Gly Leu Lys Phe Leu Asp Leu Asn Lys Asn Pro Leu Gln Arg Val Gly Pro Gly Asp Phe Ala Asn Met Leu His Leu Lys Glu Leu Gly Leu Asn Asn Met Glu Glu Leu Val Ser Ile Asp Lys Phe Ala Leu Val Asn Leu Pro Glu Leu Thr Lys Leu Asp Ile Thr Asn Asn Pro Arg Leu Ser Phe Ile His Pro Arg Ala Phe His His Leu Pro Gln Met Glu Thr Leu Met Leu Asn Asn Asn Ala Leu Ser Ala Leu His Gln Gln Thr Val Glu Ser Leu Pro Asn Leu Gln Glu Val Gly Leu His Gly Asn Pro Ile Arg Cys Asp Cys Val Ile Arg Trp Ala Asn Ala Thr Gly Thr Arg Val Arg Phe Ile Glu Pro Gln Ser Thr Leu Cys Ala Glu Pro Pro Asp Leu Gln Arg Leu Pro Val Arg Glu Val Pro Phe Arg Glu Met Thr Asp His Cys Leu Pro Leu Ile Ser Pro Arg Ser Phe Pro Pro Ser Leu Gln Val 

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Pro	Glu	Pro	Glu	Ile 455	Tyr	Trp	Val	Thr	Pro 460	Ala	Gly	Leu	Arg	Leu 465
Thr	Pro	Ala	His	Ala 470	Gly	Arg	Arg	Tyr	Arg 475	Val	Tyr	Pro	Glu	Gly 480
Thr	Leu	Glu	Leu	Arg 485	Arg	Val	Thr	Ala	Glu 490	Glu	Ala	Gly	Leu	Tyr 495
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Ser	Val	Val	Val	Gly 515	Arg	Ala	Leu	Leu	Gln 520	Pro	Gly	Arg	Asp	Glu 525
Gly	Gln	Gly	Leu	Glu 530	Leu	Arg	Val	Gln	Glu 535	Thr	His	Pro	Tyr	His 540
Ile	Leu	Leu	Ser	Trp 545	Val	Thr	Pro	Pro	Asn 550	Thr	Val	Ser	Thr	Asn 555
Leu	Thr	Trp	Ser	Ser 560	Ala	Ser	Ser	Leu	Arg 565	Gly	Gln	Gly	Ala	Thr 570
Ala	Leu	Ala	Arg	Leu 575	Pro	Arg	Gly	Thr	His 580	Ser	Tyr	Asn	Ile	Thr 585
Arg	Leu	Leu	Gln	Ala 590	Thr	Glu	Tyr	Trp	Ala 595	Cys	Leu	Gln	Val	Ala 600
Phe	Ala	Asp	Ala	His 605	Thr	Gln	Leu	Ala	Cys 610	Val	Trp	Ala	Arg	Thr 615
Lys	Glu	Ala	Thr	Ser 620	Cys	His	Arg	Ala	Leu 625	Gly	Asp	Arg	Pro	Gly 630
Leu	Ile	Ala	Ile	Leu 635	Ala	Leu	Ala	Val	Leu 640	Leu	Leu	Ala	Ala	Gly 645
Leu	Ala	Ala	His	Leu 650	Gly	Thr	Gly	Gln	Pro 655	Arg	Lys	Gly	Val	Gly 660
Gly	Arg	Arg	Pro	Leu 665	Pro	Pro	Ala	Trp	Ala 670	Phe	Trp	Gly	Trp	Ser 675
Ala	Pro	Ser	Val	Arg 680	Val	Val	Ser	Ala	Pro 685	Leu	Val	Leu	Pro	Trp 690
Asn	Pro	Gly	Arg	Lys 695	Leu	Pro	Arg	Ser	Ser 700	Glu	Gly	Glu	Thr	Leu 705
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Cys Ala His Pro Leu Ala Thr Leu Phe Lys Ile Leu Ala Ser Phe 50 55 60

Tyr Ile Ser Leu Val Ile Phe Tyr Gly Leu Ile Cys Met Tyr Thr
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Leu Trp Trp Met Leu Arg Arg Ser Leu Lys Lys Tyr Ser Phe Glu 80 85 90

Ser Ile Arg Glu Glu Ser Ser Tyr Ser Asp Ile Pro Asp Val Lys 95 100 105

Asn Asp Phe Ala Phe Met Leu His Leu Ile Asp Gln Tyr Asp Pro 110 115 120

Leu Tyr Ser Lys Arg Phe Ala Val Phe Leu Ser Glu Val Ser Glu
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Asn Lys Leu Arg Gln Leu Asn Leu Asn Asn Glu Trp Thr Leu Asp 140 145 150

Lys Leu Arg Gln Arg Leu Thr Lys Asn Ala Gln Asp Lys Leu Glu 155 160 165

Leu His Leu Phe Met Leu Ser Gly Ile Pro Asp Thr Val Phe Asp 170 175 180

Leu Val Glu Leu Glu Val Leu Lys Leu Glu Leu Ile Pro Asp Val 185 190 195

Thr Ile Pro Pro Ser Ile Ala Gln Leu Thr Gly Leu Lys Glu Leu 200 205 210

Trp Leu Tyr His Thr Ala Ala Lys Ile Glu Ala Pro Ala Leu Ala

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Ile	Thr	Ala	Asn	Arg 455	Ile	Glu	Thr	Leu	Pro 460	Pro	Glu	Leu	Phe	Gln 465
Cys	Arg	Lys	Leu	Arg 470	Ala	Leu	His	Leu	Gly 475	Asn	Asn	Val	Leu	Gln 480
Ser	Leu	Pro	Ser	Arg 485	Val	Gly	Glu	Leu	Thr 490	Asn	Leu	Thr	Gln	Ile 495

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Glu Leu Arg Gly Asn Arg Leu Glu Cys Leu Pro Val Glu Leu Gly 505 Glu Cys Pro Leu Leu Lys Arg Ser Gly Leu Val Val Glu Glu Asp Leu Phe Asn Thr Leu Pro Pro Glu Val Lys Glu Arg Leu Trp Arg 530 535 Ala Asp Lys Glu Gln Ala <210> 251 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 251 caacaatgag ggcaccaagc 20 <210> 252 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 252 gatggctagg ttctggaggt tctg 24 <210> 253 <211> 47 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 253 caacctgcag gagattgacc tcaaggacaa caacctcaag accatcg 47 <210> 254 <211> 1650 <212> DNA <213> Homo Sapien <400> 254 gcctgttgct gatgctgccg tgcggtactt gtcatggagc tggcactgcg 50 gcgctctccc gtcccgcggt ggttgctgct gctgccgctg ctgctgggcc 100 tgaacgcagg agctgtcatt gactggccca cagaggaggg caaggaagta 150

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Ala	Met	Val	Ala	Ser 140	Asp	Met	Met	Val	Leu 145	Leu	Lys	Thr	Phe	Phe 150
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Tyr	Lys	Ala	Ile	Gln 185	Arg	Gly	Thr	Ile	Lys 190	Cys	Asn	Phe	Ala	Gly 195
Val	Ala	Leu	Gly	Asp 200	Ser	Trp	Ile	Ser	Pro 205	Val	Asp	Ser	Val	Leu 210
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Ala G	Slu	Met	Ile	Ile 260	Glu	Gln	Asn	Thr	Asp 265	Gly	Val	Asn	Phe	Tyr 270
Asn I	le	Leu	Thr	Lys 275	Ser	Thr	Pro	Thr	Ser 280	Thr	Met	Glu	Ser	Ser 285
Leu G	Slu	Phe	Thr	Gln 290	Ser	His	Leu	Val	Cys 295	Leu	Cys	Gln	Arg	His 300
Val A	lrg	His	Leu	Gln 305	Arg	Asp	Ala	Leu	Ser 310	Gln	Leu	Met	Asn	Gly 315
Pro I	le	Arg	Lys	Lys 320	Leu	Lys	Ile	Ile	Pro 325	Glu	Asp	Gln	Ser	Trp 330
Gly G	Sly	Gln	Ala	Thr 335	Asn	Val	Phe	Val	Asn 340	Met	Glu	Glu	Asp	Phe 345
Met L	ys	Pro	Val	Ile 350	Ser	Ile	Val	Asp	Glu 355	Leu	Leu	Glu	Ala	Gly 360
Ile A	Asn	Val	Thr	Val 365	Tyr	Asn	Gly	Gln	Leu 370	Asp	Leu	Ile	Val	Asp 375
Thr M	let	Gly	Gln	Glu 380	Ala	Trp	Val	Arg	Lys 385	Leu	Lys	Trp	Pro	Glu 390
Leu F	?ro	Lys	Phe	Ser 395	Gln	Leu	Lys	Trp	Lys 400	Ala	Leu	Tyr	Ser	Asp 405
Pro L	Lys	Ser	Leu	Glu 410	Thr	Ser	Ala	Phe	Val 415	Lys	Ser	Tyr	Lys	Asn 420
Leu A	Ala	Phe	Tyr	Trp 425	Ile	Leu	Lys	Ala	Gly 430	His	Met	Val	Pro	Ser 435
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Gln Glu

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<212> DNA

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35 40 45

Glu Asp Ala Glu Leu Gly Arg Trp Pro Trp Gln Gly Ser Leu Arg
50 55 60

Leu Trp Asp Ser His Val Cys Gly Val Ser Leu Leu Ser His Arg

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<sup>&</sup>lt;211> 314

<sup>&</sup>lt;212> PRT

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Met	Pro	Ser	Phe	Trp 110	Ser	Leu	Gln	Ala	Tyr 115	Tyr	Thr	Arg	Tyr	Phe 120
Val	Ser	Asn	Ile	Tyr 125	Leu	Ser	Pro	Arg	Tyr 130	Leu	Gly	Asn	Ser	Pro 135
Tyr	Asp	Ile	Ala	Leu 140	Val	Lys	Leu	Ser	Ala 145	Pro	Val	Thr	Tyr	Thr 150
Lys	His	Ile	Gln	Pro 155	Ile	Cys	Leu	Gln	Ala 160	Ser	Thr	Phe	Glu	Phe 165
Glu	Asn	Arg	Thr	Asp 170	Cys	Trp	Val	Thr	Gly 175	Trp	Gly	Tyr	Ile	Lys 180
Glu	Asp	Glu	Ala	Leu 185	Pro	Ser	Pro	His	Thr 190	Leu	Gln	Glu	Val	Gln 195
Val	Ala	Ile	Ile	Asn 200	Asn	Ser	Met	Cys	Asn 205	His	Leu	Phe	Leu	Lys 210
Tyr	Ser	Phe	Arg	Lys 215	Asp	Ile	Phe	Gly	Asp 220	Met	Val	Cys	Ala	Gly 225
Asn	Ala	Gln	Gly	Gly 230	Lys	Asp	Ala	Cys	Phe 235	Gly	Asp	Ser	Gly	Gly 240
Pro	Leu	Ala	Суѕ	Asn 245	Lys	Asn	Gly	Leu	Trp 250	Tyr	Gln	Ile	Gly	Val 255
Val	Ser	Trp	Gly	Val 260	Gly	Cys	Gly	Arg	Pro 265	Asn	Arg	Pro	Gly	Val 270
Tyr	Thr	Asn	Ile	Ser 275	His	His	Phe	Glu	Trp 280	Ile	Gln	Lys	Leu	Met 285
Ala	Gln	Ser	Gly	Met 290	Ser	Gln	Pro	Asp	Pro 295	Ser	Trp	Pro	Leu	Leu 300
Phe	Phe	Pro	Leu	Leu 305	Trp	Ala	Leu	Pro	Leu 310	Leu	Gly	Pro	Val	

<sup>&</sup>lt;210> 258

<sup>&</sup>lt;211> 2427

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Homo Sapien

<sup>&</sup>lt;400> 258

cctcctaggg ctctttgccc tcatcctctc tggcaaatgc agttacagcc 100 cggagcccga ccagcggagg acgctgccc caggctgggt gtccctgggc 150 cgtgcggacc ctgaggaaga gctgagtctc acctttgccc tgagacagca 200 gaatgtggaa agactctcgg agctggtgca ggctgtgtcg gatcccagct 250 ctcctcaata cggaaaatac ctgaccctag agaatgtggc tgatctggtg 300 aggccatccc cactgaccct ccacacggtg caaaaatggc tcttggcagc 350 cggagcccag aagtgccatt ctgtgatcac acaggacttt ctgacttgct 400 ggctgagcat ccgacaagca gagctgctgc tccctggggc tgagtttcat 450 cactatgtgg gaggacctac ggaaacccat gttgtaaggt ccccacatcc 500 ctaccagett ccacaggeet tggeeceeca tgtggaettt gtggggggae 550 tgcaccgttt tcccccaaca tcatccctga ggcaacgtcc tgagccgcag 600 gtgacaggga ctgtaggcct gcatctgggg gtaaccccct ctgtgatccg 650 taagcgatac aacttgacct cacaagacgt gggctctggc accagcaata 700 acagccaagc ctgtgcccag ttcctggagc agtatttcca tgactcagac 750 ctggctcagt tcatgcgcct cttcggtggc aactttgcac atcaggcatc 800 ccagtctaga tgtgcagtac ctgatgagtg ctggtgccaa catctccacc 900 tgggtctaca gtagccctgg ccggcatgag ggacaggagc ccttcctgca 950 gtggctcatg ctgctcagta atgagtcagc cctgccacat gtgcatactg 1000 tgagctatgg agatgatgag gactccctca gcagcgccta catccagcgg 1050 gtcaacactg agctcatgaa ggctgccgct cggggtctca ccctgctctt 1100 cgcctcaggt gacagtgggg ccgggtgttg gtctgtctct ggaagacacc 1150 agttccgccc taccttccct gcctccagcc cctatgtcac cacagtggga 1200 ggcacatect tecaggaace ttteeteate acaaatgaaa ttgttgaeta 1250 tatcagtggt ggtggcttca gcaatgtgtt cccacggcct tcataccagg 1300 aggaagetgt aacgaagtte etgageteta geceecacet gecaceatee 1350 agttacttca atgccagtgg ccgtgcctac ccagatgtgg ctgcactttc 1400 tgatggctac tgggtggtca gcaacagagt gcccattcca tgggtgtccg 1450

gaacctcggc ctctactcca gtgtttgggg ggatcctatc cttgatcaat 1500 gagcacagga teettagtgg eegeeeect ettggettte teaacceaag 1550 getetaceag cageatgggg caggtetett tgatgtaace cgtggetgee 1600 atgagtcctg tctggatgaa gaggtagagg gccagggttt ctgctctggt 1650 cctggctggg atcctgtaac aggctgggga acaccaactt cccagctttg 1700 ctgaagactc tactcaaccc ctgacccttt cctatcagga gagatggctt 1750 gtcccctgcc ctgaagctgg cagttcagtc ccttattctg ccctgttgga 1800 agecetgetg aacceteaac tattgactge tgeagacage ttateteeet 1850 aaccctgaaa tgctgtgagc ttgacttgac tcccaaccct accatgctcc 1900 atcatactca ggtctcccta ctcctgcctt agattcctca ataagatgct 1950 gtaactagca ttttttgaat gcctctccct ccgcatctca tctttctctt 2000 ttcaatcagg cttttccaaa gggttgtata cagactctgt gcactatttc 2050 acttgatatt cattccccaa ttcactgcaa ggagacctct actgtcaccg 2100 tttactcttt cctaccctga catccagaaa caatggcctc cagtgcatac 2150 ttctcaatct ttgctttatg gcctttccat catagttgcc cactccctct 2200 cettaettag ettecaggte ttaacttete tgaetaetet tgtetteete 2250 tctcatcaat ttctgcttct tcatggaatg ctgaccttca ttgctccatt 2300 tgtagatttt tgctcttctc agtttactca ttgtcccctg gaacaaatca 2350 ctgacatcta caaccattac catctcacta aataagactt tctatccaat 2400 aatgattgat acctcaaatg taaaaaa 2427

- <210> 259
- <211> 556
- <212> PRT
- <213> Homo Sapien

## <400> 259

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Ser Gly Lys Cys Ser Tyr Ser Pro Glu Pro Asp Gln Arg Arg Thr 20 25 30

Leu Pro Pro Gly Trp Val Ser Leu Gly Arg Ala Asp Pro Glu Glu 35 40 45

Glu Leu Ser Leu Thr Phe Ala Leu Arg Gln Gln Asn Val Glu Arg
50 55 60

Leu	. Ser	Glu	Leu	Val	. Gln	Ala	a Val	Ser	Asp 70		Ser	Ser	Pro	Gln 75
Tyr	Gly	Lys	Туг	Leu 80	Thr	Leu	ı Glu	a Asn	Val 85		a Asp	Leu	ı Val	Arg
Pro	Ser	Pro	Leu	Thr 95	Leu	His	Thr	Val	Gln 100		Trp	Leu	Leu	Ala 105
Ala	Gly	Ala	Gln	Lys	Cys	His	Ser	· Val	Ile 115		Gln	Asp	Phe	Leu 120
Thr	Cys	Trp	Leu	Ser 125	Ile	Arg	Gln	Ala	Glu 130		Leu	Leu	Pro	Gly 135
Ala	Glu	Phe	His	His 140	Tyr	Val	Gly	Gly	Pro 145	Thr	Glu	Thr	His	Val 150
Val	Arg	Ser	Pro	His 155	Pro	Tyr	Gln	Leu	Pro 160	Gln	Ala	Leu	Ala	Pro 165
His	Val	Asp	Phe	Val 170	Gly	Gly	Leu	His	Arg 175	Phe	Pro	Pro	Thr	Ser 180
Ser	Leu	Arg	Gln	Arg 185	Pro	Glu	Pro	Gln	Val 190	Thr	Gly	Thr	Val	Gly 195
Leu	His	Leu	Gly	Val 200	Thr	Pro	Ser	Val	Ile 205	Arg	Lys	Arg	Tyr	Asn 210
Leu	Thr	Ser	Gln	Asp 215	Val	Gly	Ser	Gly	Thr 220	Ser	Asn	Asn	Ser	Gln 225
Ala	Суз	Ala	Gln	Phe 230	Leu	Glu	Gln	Tyr	Phe 235	His	Asp	Ser	Asp	Leu 240
Ala	Gln	Phe	Met	Arg 245	Leu	Phe	Gly	Gly	Asn 250	Phe	Ala	His	Gln	Ala 255
Ser	Val	Ala	Arg	Val 260	Val	Gly	Gln	Gln	Gly 265	Arg	Gly	Arg	Ala	Gly 270
Ile	Glu	Ala	Ser	Leu 275	Asp	Val	Gln	Tyr	Leu 280	Met	Ser	Ala	Gly	Ala 285
Asn	Ile	Ser	Thr	Trp 290	Val	Tyr	Ser	Ser	Pro 295	Gly	Arg	His	Glu	Gly 300
Gln	Glu	Pro	Phe	Leu 305	Gln	Trp	Leu	Met	Leu 310	Leu	Ser	Asn	Glu	Ser 315
Ala	Leu	Pro	His	Val 320	His	Thr	Val	Ser	Tyr 325	Gly	Asp	Asp	Glu	Asp 330
Ser	Leu	Ser	Ser	Ala 335	Tyr	Ile	Gln	Arg	Val 340	Asn	Thr	Glu	Leu	Met 345

Lys	Ala	Ala	Ala	Arg 350	Gly	Leu	Thr	Leu	Leu 355	Phe	Ala	Ser	Gly	Asp 360
Ser	Gly	Ala	Gly	Cys 365	Trp	Ser	Val	Ser	Gly 370	Arg	His	Gln	Phe	Arg 375
Pro	Thr	Phe	Pro	Ala 380	Ser	Ser	Pro	Tyr	Val 385	Thr	Thr	Val	Gly	Gly 390
Thr	Ser	Phe	Gln	Glu 395	Pro	Phe	Leu	Ile	Thr 400	Asn	Glu	Ile	Val	Asp 405
Tyr	Ile	Ser	Gly	Gly 410	Gly	Phe	Ser	Asn	Val 415	Phe	Pro	Arg	Pro	Ser 420
Tyr	Gln	Glu	Glu	Ala 425	Val	Thr	Lys	Phe	Leu 430	Ser	Ser	Ser	Pro	His 435
Leu	Pro	Pro	Ser	Ser 440	Tyr	Phe	Asn	Ala	Ser 445	Gly	Arg	Ala	Tyr	Pro 450
Asp	Val	Ala	Ala	Leu 455	Ser	Asp	Gly	Tyr	Trp 460	Val	Val	Ser	Asn	Arg 465
Val	Pro	Ile	Pro	Trp 470	Val	Ser	Gly	Thr	Ser 475	Ala	Ser	Thr	Pro	Val 480
Phe	Gly	Gly	Ile	Leu 485	Ser	Leu	Ile	Asn	Glu 490	His	Arg	Ile	Leu	Ser 495
Gly	Arg	Pro	Pro	Leu 500	Gly	Phe	Leu	Asn	Pro 505	Arg	Leu	Tyr	Gln	Gln 510
His	Gly	Ala	Gly	Leu 515	Phe	Asp	Val	Thr	Arg 520	Gly	Cys	His	Glu	Ser 525
Cys	Leu	Asp	Glu	Glu 530	Val	Glu	Gly	Gln	Gly 535	Phe	Cys	Ser	Gly	Pro 540
Gly	Trp	Asp	Pro	Val 545	Thr	Gly	Trp	Gly	Thr 550	Pro	Thr	Ser	Gln	Leu 555
Cys														

<210> 260 <211> 1638

<212> DNA

<213> Homo Sapien

<400> 260

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attccagggc teetetteet tetettett etgetetgtg etgttgggca 150

agtgagccct tacagtgccc cctggaaacc cacttggcct gcataccgcc 200 tecetgicgi ettgeceeag tetaceetea atttageeaa geeagaetti 250 ggagccgaag ccaaattaga agtatcttct tcatgtggac cccagtgtca 300 taagggaact ccactgccca cttacgaaga ggccaagcaa tatctgtctt 350 atgaaacgct ctatgccaat ggcagccgca cagagacgca ggtgggcatc 400 tacatcctca gcagtagtgg agatggggcc caacaccgag actcagggtc 450 ttcaggaaag tctcgaagga agcggcagat ttatggctat gacagcaggt 500 tcagcatttt tgggaaggac ttcctgctca actacccttt ctcaacatca 550 gtgaagttat ccacgggctg caccggcacc ctggtggcag agaagcatgt 600 cctcacaget geceaetgea tacaegatgg aaaaacetat gtgaaaggaa 650 cccagaaget tegagtggge tteetaaage ecaagtttaa agatggtggt 700 cgaggggcca acgactccac ttcagccatg cccgagcaga tgaaatttca 750 gtggatccgg gtgaaacgca cccatgtgcc caagggttgg atcaagggca 800 atgccaatga catcggcatg gattatgatt atgccctcct ggaactcaaa 850 aagccccaca agagaaaatt tatgaagatt ggggtgagcc ctcctgctaa 900 gcagctgcca gggggcagaa ttcacttctc tggttatgac aatgaccgac 950 caggcaattt ggtgtatcgc ttctgtgacg tcaaagacga gacctatgac 1000 ttgctctacc agcaatgcga tgcccagcca ggggccagcg ggtctggggt 1050 ctatgtgagg atgtggaaga gacagcagca gaagtgggag cgaaaaatta 1100 ttggcatttt ttcagggcac cagtgggtgg acatgaatgg ttccccacag 1150 gatttcaacg tggctgtcag aatcactcct ctcaaatatg cccagatttg 1200 ctattggatt aaaggaaact acctggattg tagggagggg tgacacagtg 1250 ttccctcctg gcagcaatta agggtcttca tgttcttatt ttaggagagg 1300 ccaaattgtt ttttgtcatt ggcgtgcaca cgtgtgtgtg tgtgtgtgt 1350 tgtgtgtaag gtgtcttata atcttttacc tatttcttac aattgcaaga 1400 tgactggctt tactatttga aaactggttt gtgtatcata tcatatatca 1450 tttaagcagt ttgaaggcat acttttgcat agaaataaaa aaaatactga 1500 tttggggcaa tgaggaatat ttgacaatta agttaatctt cacgtttttg 1550 caaactttga tttttatttc atctgaactt gtttcaaaga tttatattaa 1600

## atatttggca tacaagagat atgaaaaaaa aaaaaaaa 1638

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Cys	Ala	Val	Gly	Gln 20	Val	Ser	Pro	Tyr	Ser 25	Ala	Pro	Trp	Lys	Pro 30
Thr	Trp	Pro	Ala	Tyr 35		Leu	Pro	Val	Val 40	Leu	Pro	Gln	Ser	Thr 45
Leu	Asn	Leu	Ala	Lys 50	Pro	Asp	Phe	Gly	Ala 55	Glu	Ala	Lys	Leu	Glu 60
Val	Ser	Ser	Ser	Cys 65	Gly	Pro	Gln	Cys	His 70	Lys	Gly	Thr	Pro	Leu 75
Pro	Thr	Tyr	Glu	Glu 80	Ala	Lys	Gln	Tyr	Leu 85	Ser	Tyr	Glu	Thr	Leu 90
Tyr	Ala	Asn	Gly		Arg	Thr	Glu	Thr	Gln	Val	Gly	Ile	Tyr	Ile
				95					100					105
Leu	Ser	Ser	Ser	Gly 110	Asp	Gly	Ala	Gln	His 115	Arg	Asp	Ser	Gly	Ser 120
Ser	Gly	Lys	Ser	Arg 125	Arg	Lys	Arg	Gln	Ile 130	Tyr	Gly	Tyr	Asp	Ser 135
Arg	Phe	Ser	Ile	Phe 140	Gly	Lys	Asp	Phe	Leu 145	Leu	Asn	Tyr	Pro	Phe 150
Ser	Thr	Ser	Val	Lys 155	Leu	Ser	Thr	Gly	Cys 160	Thr	Gly	Thr	Leu	Val 165
Ala	Glu	Lys	His	Val 170	Leu	Thr	Ala	Ala	His 175	Cys	Ile	His	Asp	Gly 180
Lys	Thr	Tyr	Val	Lys 185	Gly	Thr	Gln	Lys	Leu 190	Arg	Val	Gly	Phe	Leu 195
Lys	Pro	Lys	Phe	Lys 200	Asp	Gly	Gly	Arg	Gly 205	Ala	Asn	Asp	Ser	Thr 210
Ser	Ala	Met	Pro	Glu 215	Gln	Met	Lys		Gln 220	Trp	Ile	Arg	Val	Lys 225
Arg	Thr	His	Val	Pro 230	Lys	Gly	Trp		Lys 235	Gly	Asn	Ala	Asn	Asp 240

Ile Gly Met Asp Tyr Asp Tyr Ala Leu Leu Glu Leu Lys Lys Pro 245 His Lys Arg Lys Phe Met Lys Ile Gly Val Ser Pro Pro Ala Lys 260 265 Gln Leu Pro Gly Gly Arg Ile His Phe Ser Gly Tyr Asp Asn Asp Arg Pro Gly Asn Leu Val Tyr Arg Phe Cys Asp Val Lys Asp Glu 290 Thr Tyr Asp Leu Leu Tyr Gln Gln Cys Asp Ala Gln Pro Gly Ala Ser Gly Ser Gly Val Tyr Val Arg Met Trp Lys Arg Gln Gln 320 Lys Trp Glu Arg Lys Ile Ile Gly Ile Phe Ser Gly His Gln Trp Val Asp Met Asn Gly Ser Pro Gln Asp Phe Asn Val Ala Val Arg 355 350 Ile Thr Pro Leu Lys Tyr Ala Gln Ile Cys Tyr Trp Ile Lys Gly 370 365

Asn Tyr Leu Asp Cys Arg Glu Gly 380

<210> 262

<211> 1378

<212> DNA

<213> Homo Sapien

<400> 262

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<210> 263
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<400> 263

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Asn Ala Ala Arg Ile Pro Val Pro Pro Ala Cys Gly Lys Pro Gln 35 40 45

Gln Leu Asn Arg Val Val Gly Gly Glu Asp Ser Thr Asp Ser Glu
50 55 60

Trp Pro Trp Ile Val Ser Ile Gln Lys Asn Gly Thr His His Cys
65 70 75

<sup>&</sup>lt;211> 317

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo Sapien

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Cys	Phe	Lys	Asp	Asn 95	Leu	Asn	Lys	Pro	Tyr 100	Leu	Phe	Ser	Val	Leu 105
Leu	Gly	Ala	Trp	Gln 110	Leu	Gly	Asn	Pro	Gly 115	Ser	Arg	Ser	Gln	Lys 120
Val	Gly	Val	Ala	Trp 125	Val	Glu	Pro	His	Pro 130	Val	Tyr	Ser	Trp	Lys 135
Glu	Gly	Ala	Cys	Ala 140	Asp	Ile	Ala	Leu	Val 145	Arg	Leu	Glu	Arg	Ser 150
Ile	Gln	Phe	Ser	Glu 155	Arg	Val	Leu	Pro	Ile 160	Cys	Leu	Pro	Asp	Ala 165
Ser	Ile	His	Leu	Pro 170	Pro	Asn	Thr	His	Cys 175	Trp	Ile	Ser	Gly	Trp 180
Gly	Ser	Ile	Gln	Asp 185	Gly	Val	Pro	Leu	Pro 190	His	Pro	Gln	Thr	Leu 195
Gln	Lys	Leu	Lys	Val 200	Pro	Ile	Ile	Asp	Ser 205	Glu	Val	Cys	Ser	His 210
Leu	Tyr	Trp	Arg	Gly 215	Ala	Gly	Gln	Gly	Pro 220	Ile	Thr	Glu	Asp	Met 225
Leu	Cys	Ala	Gly	Tyr 230	Leu	Glu	Gly	Glu	Arg 235	Asp	Ala	Cys	Leu	Gly 240
Asp	Ser	Gly	Gly	Pro 245	Leu	Met	Cys	Gln	Val 250	Asp	Gly	Ala	Trp	Leu 255
Leu	Ala	Gly	Ile	Ile 260	Ser	Trp	Gly	Glu	Gly 265	Суѕ	Ala	Glu	Arg	Asn 270
Arg	Pro	Gly	Val	Tyr 275	Ile	Ser	Leu	Ser	Ala 280	His	Arg	Ser	Trp	Val 285
Glu	Lys	Ile	Val	Gln 290	Gly	Val	Gln	Leu	Arg 295	Gly	Arg	Ala	Gln	Gly 300
Gly	Gly	Ala	Leu	Arg 305	Ala	Pro	Ser	Gln	Gly 310	Ser	Gly	Ala	Ala	Ala 315

Arg Ser

<210> 264 <211> 24

<212> DNA

<213> Artificial Sequence

<220>

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<210> 265
<211> 19
<212> DNA
<213> Artificial Sequence
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<223> Synthetic Oligonucleotide Probe
<400> 265
gcagaggtgt ctaaggttg 19
<210> 266
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 266
agetetagae caatgeeage ttee 24
<210> 267
<211> 45
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 267
qccaccaact cctgcaagaa cttctcagaa ctgcccctgg tcatg 45
<210> 268
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
ggggaattca ccctatgaca ttgcc 25
<210> 269
<211> 24
<212> DNA
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<220>
<223> Synthetic Oligonucleotide Probe
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<210> 271
<211> 26
<212> DNA
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<223> Synthetic Oligonucleotide Probe
<400> 271
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<210> 272
<211> 18
<212> DNA
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<223> Synthetic Oligonucleotide Probe
<400> 272
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 <400> 273
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 <210> 274
 <211> 24
 <212> DNA
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 <400> 274
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<211> 45
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
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<211> 21
<212> DNA
<213> Artificial Sequence
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<400> 276
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<210> 277
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 277
 ggctatgaca gcaggttc 18
<210> 278
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 278
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<210> 279
<211> 24
<212> DNA
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<212> DNA
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<223> Synthetic Oligonucleotide Probe
<400> 281
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<210> 282
<211> 61
<212> DNA
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<220>
<223> Synthetic Oligonucleotide Probe
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 tgccaggtgg a 61
<210> 283
<211> 119
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
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<212> PRT

<213> Homo Sapien

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Glu Glu Lys Arg Leu Met Val Glu Leu His Asn Leu Tyr Arg Ala 35 40 45

Gln Val Ser Pro Thr Ala Ser Asp Met Leu His Met Arg Trp Asp
50 55 60

Glu Glu Leu Ala Ala Phe Ala Lys Ala Tyr Ala Arg Gln Cys Val 65 70 75

Trp Gly His Asn Lys Glu Arg Gly Arg Arg Gly Glu Asn Leu Phe
80 85 90

Ala Ile Thr Asp Glu Gly Met Asp Val Pro Leu Ala Met Glu Glu
95 100 105

Trp His His Glu Arg Glu His Tyr Asn Leu Ser Ala Ala Thr Cys 110 115 120

Ser Pro Gly Gln Met Cys Gly His Tyr Thr Gln Val Val Trp Ala 125 130 135

Lys Thr Glu Arg Ile Gly Cys Gly Ser His Phe Cys Glu Lys Leu 140 145 150

Gln Gly Val Glu Glu Thr Asn Ile Glu Leu Leu Val Cys Asn Tyr 155 160 165

Glu Pr	o Pro	Gly	Asn 170	Val	Lys	Gly	Lys	Arg 175	Pro	Tyr	Gln	Glu	Gly 180
Thr Pr	o Cys	Ser	Gln 185	Cys	Pro	Ser	Gly	Tyr 190	His	Cys	Lys	Asn	Ser 195
Leu Cy	s Glu	Pro	Ile 200	Gly	Ser	Pro	Glu	Asp 205	Ala	Gln	Asp	Leu	Pro 210
Tyr Le	eu Val	Thr	Glu 215	Ala	Pro	Ser	Phe	Arg 220	Ala	Thr	Glu	Ala	Ser 225
Asp Se	er Arg	Lys	Met 230	Gly	Thr	Pro	Ser	Ser 235	Leu	Ala	Thr	Gly	Ile 240
Pro Al	a Phe	Leu	Val 245	Thr	Glu	Val	Ser	Gly 250	Ser	Leu	Ala	Thr	Lys 255
Ala Le	eu Pro	Ala	Val 260	Glu	Thr	Gln	Ala	Pro 265	Thr	Ser	Leu	Ala	Thr 270
Lys As	sp Pro	Pro	Ser 275	Met	Ala	Thr	Glu	Ala 280	Pro	Pro	Cys	Val	Thr 285
Thr Gl	u Val	Pro	Ser 290	Ile	Leu	Ala	Ala	His 295	Ser	Leu	Pro	Ser	Leu 300
Asp Gl	u Glu	Pro	Val 305	Thr	Phe	Pro	Lys	Ser 310	Thr	His	Val	Pro	Ile 315
Pro Ly	s Ser	Ala	Asp 320	Lys	Val	Thr	Asp	Lys 325	Thr	Lys	Val	Pro	Ser 330
Arg Se	er Pro	Glu	Asn 335	Ser	Leu	Asp	Pro	Lys 340	Met	Ser	Leu	Thr	Gly 345
Ala Ar	g Glu	Leu	Leu 350	Pro	His	Ala	Gln	Glu 355	Glu	Ala	Glu	Ala	Glu 360
Ala Gl	u Leu	Pro	Pro 365	Ser	Ser	Glu	Val	Leu 370	Ala	Ser	Val	Phe	Pro 375
Ala Gl	n Asp	Lys	Pro 380	Gly	Glu	Leu	Gln	Ala 385	Thr	Leu	Asp	His	Thr 390
Gly Hi	s Thr	Ser	Ser 395	Lys	Ser	Leu	Pro	Asn 400	Phe	Pro	Asn	Thr	Ser 405
Ala Th	r Ala	Asn	Ala 410	Thr	Gly	Gly	Arg	Ala 415	Leu	Ala	Leu	Gln	Ser 420
Ser Le	u Pro	Gly	Ala 425	Glu	Gly	Pro	Asp	Lys 430	Pro	Ser	Val	Val	Ser 435
Gly Le	u Asn	Ser	Gly 440	Pro	Gly	His	Val	Trp 445	Gly	Pro	Leu	Leu	Gly 450

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35 40 45

Asn Asn Glu Leu Glu Thr Ile Pro Asn Leu Gly Pro Val Ser 50 55 60

Ala Asn Ile Thr Leu Leu Ser Leu Ala Gly Asn Arg Ile Val Glu 65 70 75

Ile Leu Pro Glu His Leu Lys Glu Phe Gln Ser Leu Glu Thr Leu 80 85 90

Asp Leu Ser Ser Asn Asn Ile Ser Glu Leu Gln Thr Ala Phe Pro 95 100 105

Ala Leu Gln Leu Lys Tyr Leu Tyr Leu Asn Ser Asn Arg Val Thr 110 115 120

Ser Met Glu Pro Gly Tyr Phe Asp Asn Leu Ala Asn Thr Leu Leu 125 130 135

Val Leu Lys Leu Asn Arg Asn Arg Ile Ser Ala Ile Pro Pro Lys 140 145 150

Met Phe Lys Leu Pro Gln Leu Gln His Leu Glu Leu Asn Arg Asn

Lys Ile Lys Asn Val Asp Gly Leu Thr Phe Gln Gly Leu Gly All Reu Lys Ser Leu Lys Met Gln Arg Asn Gly Val Thr Lys Leu Me 185															•
Leu Lys Ser Leu Lys Met Gln Arg Asn Gly Val Thr Lys Leu Me 185  Asp Gly Ala Phe Trp Gly Leu Ser Asn Met Glu Ile Leu Gln Le 200  Asp His Asn Asn Leu Thr Glu Ile Thr Lys Gly Trp Leu Tyr Gl 222  Leu Leu Met Leu Gln Glu Leu His Leu Ser Gln Asn Ala Ile As 235  Arg Ile Ser Pro Asp Ala Trp Glu Phe Cys Gln Lys Leu Ser Gl 255  Leu Asp Leu Thr Phe Asn His Leu Ser Arg Leu Asp Asp Ser Se 265  Leu Gly Leu Ser Leu Leu Asn Thr Leu His Ile Gly Asn As 288  Arg Val Ser Tyr Ile Ala Asp Cys Ala Phe Arg Gly Leu Ser Se 295  Leu Lys Thr Leu Asp Leu Lys Asn Asn Glu Ile Ser Trp Thr Il 305  Glu Asp Met Asn Gly Ala Phe Ser Gly Leu Asp Lys Leu Arg Arg 336  Leu Ile Leu Gln Gly Asn Arg Ile Arg Ser Ile Thr Lys Lys Al 336  Ala Ile Met Ser Leu Gln Gly Asn And Phe Ser Gly Leu Asp Leu Ser Asp Asp And Ala Ile Met Lys Trp Leu Asp And Thr Ser Ser Leu Leu Asn Thr Leu Asp Leu Cys Asp And And Ile Cys Asp Cyn And And Ile Cys And And And And Ile Cys And And And Ile Cys And And And And Ile Cys And And And And Ile Cys And					155					160					165
Asp Gly Ala Phe Trp Gly Leu Ser Asn Met Glu Ile Leu Gln Le 200	Lys	Ile	Lys	Asn		Asp	Gly	Leu	Thr		Gln	Gly	Leu	Gly	Ala 180
200 205 21  Asp His Asn Asn Leu Thr Glu Ile Thr Lys Gly Trp Leu Tyr Gl 222  Leu Leu Met Leu Gln Glu Leu His Leu Ser Gln Asn Ala Ile As 230  Arg Ile Ser Pro Asp Ala Trp Glu Phe Cys Gln Lys Leu Ser Gl 255  Leu Asp Leu Thr Phe Asn His Leu Ser Arg Leu Asp Asp Ser Se 266  Phe Leu Gly Leu Ser Leu Leu Asn Thr Leu His Ile Gly Asn As 288  Arg Val Ser Tyr Ile Ala Asp Cys Ala Phe 295  Leu Lys Thr Leu Asp Leu Lys Asn Asn Glu Ile Ser Trp Thr Il 305  Glu Asp Met Asn Gly Ala Phe Ser Gly Leu Asp Lys Leu Arg Ar 325  Leu Ile Leu Gln Gly Asn Arg Ile Arg Ser Ile Thr Lys Lys Al 335  Ala Ile Met Ser Leu Gln Gly Asn Thr Ser Ser Leu Leu Cys Asp Asp Cys 389  Gln Leu Lys Trp Leu Asp Leu Asn Thr Ser Ser Leu Leu Cys Asp Cy 380  Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Asn Phe Gl 410  Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gl 410  Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Ada 430	Leu	Lys	Ser	Leu		Met	Gln	Arg	Asn		Val	Thr	Lys	Leu	Met 195
Leu Leu Met Leu Gln Glu Leu His Leu Ser Gln Asn Ala Ile As 230  Arg Ile Ser Pro Asp Ala Trp Glu Phe Cys Gln Lys Leu Ser Gl 245  Leu Asp Leu Thr Phe Asn His Leu Ser Arg Leu Asp Asp Ser Se 260  Phe Leu Gly Leu Ser Leu Leu Asn Thr Leu His Ile Gly Asn As 280  Arg Val Ser Tyr Ile Ala Asp Cys Ala Phe Arg Gly Leu Ser Se 295  Leu Lys Thr Leu Asp Leu Lys Asn Asn Glu Ile Ser Trp Thr Il 305  Glu Asp Met Asn Gly Ala Phe Ser Gly Leu Asp Lys Leu Arg Ar 325  Ala Ile Leu Gln Gly Leu Asp Ala Leu Glu His Leu Asp Leu Ser Asp As 355  Ala Ile Met Ser Leu Gln Gly Asn Thr Ser Ser Leu Leu Cys Asp Cys Asp Cys 366  Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gl 400  Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gl 425  Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Asp Asp Asp Asp Asp As	Asp	Gly	Ala	Phe	_	Gly	Leu	Ser	Asn		Glu	Ile	Leu	Gln	Leu 210
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Leu Asp Leu Thr       Phe Asn His Leu Ser Arg Leu Asp Asp Ser Se 265         Phe Leu Gly Leu Ser Leu Leu Leu Asn Thr Leu His Ile Gly Asn As 28         Arg Val Ser Tyr Ile Ala Asp Cys Ala Phe Arg Gly Leu Ser Se 295         Leu Lys Thr Leu Asp Leu Lys Asn Asn Glu Ile Ser Trp Thr Il 310         Glu Asp Met Asn Gly Ala Phe Ser Gly Leu Asp Lys Leu Arg Arg 320         Leu Ile Leu Gln Gly Asn Arg Ile Arg Ser Ile Thr Lys Lys Alg 335         Phe Thr Gly Leu Asp Ala Leu Glu His Leu Asp Leu Ser Asp Asg 350         Ala Ile Met Ser Leu Gln Gly Asn Ala Phe Ser Gly His Leu Asp Leu Ser Asp Asg 365         Ala Ile Met Ser Leu Gln Gly Asn Ala Phe Ser Glu His Leu Asp Leu Ser Asp Asg 365         Ala Ile Met Ser Leu Gln Gly Asn Ala Phe Ser Gln Met Lys Lyg 370         Leu Gln Gln Leu His Leu Asn Thr Ser Ser Leu Leu Cys Asp Cy 389         Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gl 400         Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gl 410         Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asg 430	Leu	Leu	Met	Leu		Glu	Leu	His	Leu		Gln	Asn	Ala	Ile	Asn 240
260	Arg	Ile	Ser	Pro	_	Ala	Trp	Glu	Phe		Gln	Lys	Leu	Ser	Glu 255
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Leu Lys Thr Leu Asp Leu Lys Asn Asn Glu Ile Ser Trp Thr Il 31  Glu Asp Met Asn Gly Ala Phe Ser Gly Leu Asp Lys Leu Arg Ar 325  Leu Ile Leu Gln Gly Asn Arg Ile Arg Ser Ile Thr Lys Lys Al 340  Phe Thr Gly Leu Asp Ala Leu Glu His Leu Asp Leu Ser Asp As 365  Ala Ile Met Ser Leu Gln Gly Asn Thr Ser Ser Leu Leu Cys Asp Cy 380  Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gl 40  Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gla Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Asp Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asp Asp Asp Asp Asp Asp Asp Asp As	Phe	Leu	Gly	Leu		Leu	Leu	Asn	Thr		His	Ile	Gly	Asn	Asn 285
Glu Asp Met Asn Gly Ala Phe Ser Gly Leu Asp Lys Leu Arg Arg 325  Leu Ile Leu Gln Gly Asn Arg Ile Arg Ser Ile Thr Lys Lys Algand  Phe Thr Gly Leu Asp Ala Leu Glu His Leu Asp Leu Ser Asp Asg 365  Ala Ile Met Ser Leu Gln Gly Asn Thr Ser Ser Leu Leu Cys Asp Cy 385  Cleu Gln Gln Leu His Leu Asn Thr Ser Ser Leu Leu Cys Asp Cy 385  Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gla 400  Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Glar Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp Asg	Arg	Val	Ser	Tyr		Ala	Asp	Cys	Ala		Arg	Gly	Leu	Ser	Ser 300
Leu Ile Leu Gln Gly Asn Arg Ile Arg Ser Ile Thr Lys Lys Al 349  Phe Thr Gly Leu Asp Ala Leu Glu His Leu Asp Leu Ser Asp As 355  Ala Ile Met Ser Leu Gln Gly Asn Ala Phe Ser Gln Met Lys Ly 370  Leu Gln Gln Leu His Leu Asn Thr Ser Ser Leu Leu Cys Asp Cy 385  Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gl 400  Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gl 415  Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp As As 43	Leu	Lys	Thr	Leu		Leu	Lys	Asn	Asn		Ile	Ser	Trp	Thr	Ile 315
Phe Thr Gly Leu Asp Ala Leu Glu His Leu Asp Leu Ser Asp As 36  Ala Ile Met Ser Leu Gln Gly Asn Ala Phe Ser Gln Met Lys Ly 365  Leu Gln Gln Leu His Leu Asn Thr Ser Ser Leu Leu Cys Asp Cy 380  Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gl 400  Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gl 410  Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp As 43	Glu	Asp	Met	Asn		Ala	Phe	Ser	Gly		Asp	Lys	Leu	Arg	Arg 330
Ala Ile Met Ser Leu Gln Gly Asn Ala Phe Ser Gln Met Lys Ly 377  Leu Gln Gln Leu His Leu Asn Thr Ser Ser Leu Leu Cys Asp Cy 380  Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gl 400  Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gl 415  Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp As 43	Leu	Ile	Leu	Gln	_	Asn	Arg	Ile	Arg		Ile	Thr	Lys	Lys	Ala 345
Leu Gln Gln Leu His Leu Asn Thr Ser Ser Leu Leu Cys Asp Cy 385  Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gl 400  Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gl 410  Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp As 430	Phe	Thr	Gly	Leu		Ala	Leu	Glu	His		Asp	Leu	Ser	Asp	Asn 360
Gln Leu Lys Trp Leu Pro Gln Trp Val Ala Glu Asn Asn Phe Gl 400  Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gl 410  Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp As 430	Ala	Ile	Met	Ser		Gln	Gly	Asn	Ala		Ser	Gln	Met	Lys	Lys 375
Ser Phe Val Asn Ala Ser Cys Ala His Pro Gln Leu Leu Lys Gl 410  Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp As 425	Leu	Gln	Gln	Leu		Leu	Asn	Thr	Ser		Leu	Leu	Cys	Asp	Cys 390
Arg Ser Ile Phe Ala Val Ser Pro Asp Gly Phe Val Cys Asp As 425 430 43	Gln	Leu	Lys	Trp		Pro	Gln	Trp	Val		Glu	Asn	Asn	Phe	Gln 405
425 430 43	Ser	Phe	Val	Asn		Ser	Cys	Ala	His		Gln	Leu	Leu	Lys	Gly 420
Phe Pro Lye Pro Gla Tle Thr Val Cla Pro Gla Thr Cla Com Al	Arg	Ser	Ile	Phe		Val	Ser	Pro	Asp		Phe	Val	Cys	Asp	Asp 435
Phe Pro Lys Pro Gln Ile Thr Val Gln Pro Glu Thr Gln Ser Al	Phe	Pro	Lys	Pro	Gln	Ile	Thr	Val	Gln	Pro	Glu	Thr	Gln	Ser	Ala

The Lys Gly Ser						440					445					450	
470 475 480  Leu His Asp Ala Glu Met Glu Asn Tyr Ala His Leu Arg Ala Gln 495  Gly Gly Glu Val Met Glu Tyr Thr Thr Ile Leu Arg Leu Arg Glu 500  Val Glu Phe Ala Ser Glu Gly Lys Tyr Gln Cys Val Ile Ser Asn 515  His Phe Gly Ser Ser Tyr Ser Val Lys Ala Lys Leu Thr Val Asn 540  Met Leu Pro Ser Phe Thr Lys Thr Pro Met Asp Leu Thr Ile Arg 555  Ala Gly Ala Met Ala Arg Leu Glu Cys Ala Ala Val Gly His Pro 565  Ala Pro Gln Ile Ala Trp Gln Lys Asp Gly Gly Thr Asp Phe Pro 575  Ala Ala Arg Glu Arg Arg Met His Val Met Pro Glu Asp Asp Val 590  Phe Phe Ile Val Asp Val Lys Ile Glu Asp Ile Gly Val Tyr Ser 615  Cys Thr Ala Gln Asn Ser Ala Gly Ser Ile Ser Ala Asn Ala Thr 620  Leu Thr Val Leu Glu Thr Pro Ser Phe Leu Arg Pro Leu Leu Asp 645  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 650  Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Lys Tyr 705  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Ary Val 715		Ile	Lys	Gly	Ser		Leu	Ser	Phe	Ile		Ser	Ala	Ala	Ser		
Leu His Asp Ala Glu Met Glu Asn Tyr Ala His Leu Arg Ala Gln 485  Gly Gly Glu Val Met Glu Tyr Thr Thr Ile Leu Arg Leu Arg Glu 500  Val Glu Phe Ala Ser Glu Gly Lys Tyr Gln Cys Val Ile Ser Asn 525  His Phe Gly Ser Ser Tyr Ser Val Lys Ala Lys Leu Thr Val Asn 530  Met Leu Pro Ser Phe Thr Lys Thr Pro Met Asp Leu Thr Ile Arg 545  Ala Gly Ala Met Ala Arg Leu Glu Cys Ala Ala Val Gly His Pro 560  Ala Pro Gln Ile Ala Trp Gln Lys Asp Gly Gly Thr Asp Phe Pro 570  Ala Ala Arg Glu Arg Arg Met His Val Met Pro Glu Asp Asp Val 590  Phe Phe Ile Val Asp Val Lys Ile Glu Asp Ile Gly Val Tyr Ser 605  Cys Thr Ala Gln Asn Ser Ala Gly Ser Ile Ser Ala Asn Ala Thr 620  Cys Thr Val Leu Glu Thr Pro Ser Phe Leu Arg Pro Leu Leu Asp 635  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 650  Gly Gly Ser Pro Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 665  Fro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Lys Tyr 700  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 715		Ser	Asp	Ser	Pro	Met	Thr	Phe	Ala	Trp	Lys	Lys	Asp	Asn	Glu	Leu	
Gly Gly Glu Val Met Glu Tyr Thr Thr Ile Leu Arg Leu Arg Glu 510  Val Glu Phe Ala Ser Glu Gly Lys Tyr Gln Cys Val Ile Ser Asn 525  His Phe Gly Ser Ser Tyr Ser Val Lys Ala Lys Leu Thr Val Asn 540  Met Leu Pro Ser Phe Thr Lys Thr Pro Met Asp Leu Thr Ile Arg 545  Ala Gly Ala Met Ala Arg Leu Glu Cys Ala Ala Val Gly His Pro 570  Ala Pro Gln Ile Ala Trp Gln Lys Asp Gly Gly Thr Asp Phe Pro 585  Ala Ala Arg Glu Arg Arg Met His Val Met Pro Glu Asp Asp Val 590  Phe Phe Ile Val Asp Val Lys Ile Glu Asp Ile Gly Val Tyr Ser 615  Cys Thr Ala Gln Asn Ser Ala Gly Ser Ile Ser Ala Asn Ala Thr 625  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 655  Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 680  Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 705  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 770						470					475					480	
Solution		Leu	His	Asp	Ala		Met	Glu	Asn	Tyr		His	Leu	Arg	Ala		
### Since ### Si		Gly	Gly	Glu	Val		Glu	Tyr	Thr	Thr		Leu	Arg	Leu	Arg		
Met Leu Pro Ser   Phe Thr Lys Thr Pro Met Asp Leu Thr Ile Arg 555		Val	Glu	Phe	Ala		Glu	Gly	Lys	Tyr		Cys	Val	Ile	Ser		
545 550 555  Ala Gly Ala Met Ala Arg Leu Glu Cys Ala Ala Val Gly His Pro 560  Ala Pro Gln Ile Ala Trp Gln Lys Asp Gly Gly Thr Asp Phe Pro 585  Ala Ala Arg Glu Arg Arg Met His Val Met Pro Glu Asp Asp Val 600  Phe Phe Ile Val Asp Val Lys Ile Glu Asp Ile Gly Val Tyr Ser 615  Cys Thr Ala Gln Asn Ser Ala Gly Ser Ile Ser Ala Asn Ala Thr 630  Leu Thr Val Leu Glu Thr Pro Ser Phe Leu Arg Pro Leu Leu Asp 645  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 660  Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 680  Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 705  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 720		His	Phe	Gly	Ser		Tyr	Ser	Val	Lys		Lys	Leu	Thr	Val		
Ala Pro Gln Ile Ala Trp Gln Lys Asp Gly Gly Thr Asp Phe Pro 585  Ala Ala Arg Glu Arg Arg Met His Val Met Pro Glu Asp Asp Val 600  Phe Phe Ile Val Asp Val Lys Ile Glu Asp Ile Gly Val Tyr Ser 615  Cys Thr Ala Gln Asn Ser Ala Gly Ser Ile Ser Ala Asn Ala Thr 630  Leu Thr Val Leu Glu Thr Pro Ser Phe Leu Arg Pro Leu Leu Asp 645  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 650  Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 690  Leu Leu Ile Ile Val Asp Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 715  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 720		Met	Leu	Pro	Ser		Thr	Lys	Thr	Pro		Asp	Leu	Thr	Ile	_	
575  Ala Ala Arg Glu Arg Arg Met His Val Met Pro Glu Asp Asp Val 600  Phe Phe Ile Val Asp Val Lys Ile Glu Asp Ile Gly Val Tyr 615  Cys Thr Ala Gln Asn Ser Ala Gly Ser Ile Ser Ala Asn Ala Thr 630  Leu Thr Val Leu Glu Thr Pro Ser Phe Leu Arg Pro Leu Leu Asp 645  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 660  Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 690  Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 705  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 720						560					565					570	
590 595 600  Phe Phe Ile Val Asp Val Lys Ile Glu Asp Ile Gly Val Tyr Ser 615  Cys Thr Ala Gln Asn Ser Ala Gly Ser Ile Ser Ala Asn Ala Thr 620  Leu Thr Val Leu Glu Thr Pro Ser Phe Leu Arg Pro Leu Leu Asp 645  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 655  Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 680  Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 705  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 715						575					580					585	
Cys Thr Ala Gln Asn Ser Ala Gly Ser Ile Ser Ala Asn Ala Thr 620  Leu Thr Val Leu Glu Thr Pro Ser Phe Leu Arg Pro Leu Leu Asp 645  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 655  Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 690  Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 705  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 715						590					595					600	
Leu Thr Val Leu Glu Thr Pro Ser Phe Leu Arg Pro Leu Leu Asp 645  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 650  Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 680  Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 700  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 710						605					610					615	
635 640 645  Arg Thr Val Thr Lys Gly Glu Thr Ala Val Leu Gln Cys Ile Ala 660  Gly Gly Ser Pro Pro Pro 665 Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 680  Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 700  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 715	·					620					625					630	
Gly Gly Ser Pro Pro Pro Lys Leu Asn Trp Thr Lys Asp Asp Ser 675  Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 680  Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 705  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 715						635					640					645	
Pro Leu Val Val Thr Glu Arg His Phe Phe Ala Ala Gly Asn Gln 680  Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 700  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 715						650					655					660	
Leu Leu Ile Ile Val Asp Ser Asp Val Ser Asp Ala Gly Lys Tyr 695  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 710  715  690  Leu Leu Ile Ile Val Asp Ser Asp Val Gly Asn Val 715						665					670					675	•
695 700 705  Thr Cys Glu Met Ser Asn Thr Leu Gly Thr Glu Arg Gly Asn Val 710 715 720						680					685					690	
710 715 720						695	-		_		700	-			_	705	
		Thr	Cys	Glu	Met		Asn	Thr	Leu	Gly		Glu	Arg	Gly	Asn	720	

Arg	Leu	Ser	Val	Ile 725	Pro	Thr	Pro	Thr	Cys 730	Asp	Ser	Pro	Gln	Met 735
Thr	Ala	Pro	Ser	Leu 740	Asp	Asp	Asp	Gly	Trp 745	Ala	Thr	Val	Gly	Val 750
Val	Ile	Ile	Ala	Val 755	Val	Cys	Cys	Val	Val 760	Gly	Thr	Ser	Leu	Val 765
Trp	Val	Val	Ile	Ile 770	Tyr	His	Thr	Arg	Arg 775	Arg	Asn	Glu	Asp	Cys 780
Ser	Ile	Thr	Asn	Thr	Asp	Glu	Thr	Asn	Leu	Pro	Ala	Asp	Ile	Pro
				785					790					795
Ser	Tyr	Leu	Ser	Ser 800	Gln	Gly	Thr	Leu	Ala 805	Asp	Arg	Gln	Asp	Gly 810
Tyr	Val	Ser	Ser	Glu 815	Ser	Gly	Ser	His	His 820	Gln	Phe	Val	Thr	Ser 825
Ser	Gly	Ala	Gly	Phe 830	Phe	Leu	Pro	Gln	His 835	Asp	Ser	Ser	Gly	Thr 840
Cys	His	Ile	Asp	Asn 845	Ser	Ser	Glu	Ala	Asp 850	Val	Glu	Ala	Ala	Thr 855
Asp	Leu	Phe	Leu	Cys 860	Pro	Phe	Leu	Gly	Ser 865	Thr	Gly	Pro	Met	Tyr 870
Leu	Lys	Gly	Asn	Val 875	Tyr	Gly	Ser	Asp	Pro 880	Phe	Glu	Thr	Tyr	His 885
Thr	Gly	Cys	Ser	Pro 890	Asp	Pro	Arg	Thr	Val 895	Leu	Met	Asp	His	Tyr 900
Glu	Pro	Ser	Tyr	Ile 905	Lys	Lys	Lys	Glu	Cys 910	Tyr	Pro	Cys	Ser	His 915
Pro	Ser	Glu	Glu	Ser 920	Суѕ	Glu	Arg	Ser	Phe 925	Ser	Asn	Ile	Ser	Trp 930
Pro	Ser	His	Val	Arg 935	Lys	Leu	Leu	Asn	Thr 940	Ser	Tyr	Ser	His	Asn 945
Glu	Gly	Pro	Gly	Met 950	Lys	Asn	Leu	Cys	Leu 955		Lys	Ser	Ser	Leu 960
Asp	Phe	Ser	Ala	Asn 965	Pro	Glu	Pro	Ala	Ser 970		Ala	Ser	Ser	Asn 975
Ser	Phe	Met	Gly	Thr 980		Gly	Lys	Ala	Leu 985		Arg	Pro	His	Leu 990
Asp	Ala	Tyr	Ser	Ser 995		Gly	Gln		Ser 1000		Cys	Gln	Pro	Arg 1005

Ala Phe Tyr Leu Lys Ala His Ser Ser Pro Asp Leu Asp Ser Gly 1010 1015 1020

Ser Glu Glu Asp Gly Lys Glu Arg Thr Asp Phe Gln Glu Glu Asn 1025 1030 1035

His Ile Cys Thr Phe Lys Gln Thr Leu Glu Asn Tyr Arg Thr Pro 1040 1045 1050

Asn Phe Gln Ser Tyr Asp Leu Asp Thr 1055

<210> 291

<211> 2906

<212> DNA

<213> Homo Sapien

<400> 291

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gcccttctgt gtgctcctgc agcaaccagt tcagcaaggt gatttgtgtt 1000 cggaaaaacc tgcgtgaggt tccggatggc atctccacca acacacggct 1050 gctgaacctc catgagaacc aaatccagat catcaaagtg aacagcttca 1100 agcacttgag gcacttggaa atcctacagt tgagtaggaa ccatatcaga 1150 accattgaaa ttggggcttt caatggtctg gcgaacctca acactctgga 1200 actetttgae aategtetta etaceatece gaatggaget tttgtataet 1250 tgtctaaact gaaggagctc tggttgcgaa acaaccccat tgaaagcatc 1300 cettettatg cttttaacag aatteettet ttgegeegae tagaettagg 1350 ggaattgaaa agactttcat acatctcaga aggtgccttt gaaggtctgt 1400 ccaacttgag gtatttgaac cttgccatgt gcaaccttcg ggaaatccct 1450 aacctcacac cgctcataaa actagatgag ctggatcttt ctgggaatca 1500 tttatctgcc atcaggcctg gctctttcca gggtttgatg caccttcaaa 1550 aactgtggat gatacagtcc cagattcaag tgattgaacg gaatgccttt 1600 gacaaccttc agtcactagt ggagatcaac ctggcacaca ataatctaac 1650 attactgcct catgacetet teactecett geateateta gageggatae 1700 atttacatca caaccettgg aactgtaact gtgacatact gtggctcage 1750 tggtggataa aagacatggc cccctcgaac acagcttgtt gtgcccggtg 1800 taacactcct cccaatctaa aggggaggta cattggagag ctcgaccaga 1850 attacttcac atgctatgct ccggtgattg tggagccccc tgcagacctc 1900 aatgtcactg aaggcatggc agctgagctg aaatgtcggg cctccacatc 1950 cctgacatct gtatcttgga ttactccaaa tggaacagtc atgacacatg 2000 gggcgtacaa agtgcggata gctgtgctca gtgatggtac gttaaatttc 2050 acaaatgtaa ctgtgcaaga tacaggcatg tacacatgta tggtgagtaa 2100 ttccgttggg aatactactg cttcagccac cctgaatgtt actgcagcaa 2150 ccactactcc tttctcttac ttttcaaccg tcacagtaga gactatggaa 2200 ccgtctcagg atgaggcacg gaccacagat aacaatgtgg gtcccactcc 2250 agtggtcgac tgggagacca ccaatgtgac cacctctctc acaccacaga 2300 gcacaaggtc gacagagaaa accttcacca tcccagtgac tgatataaac 2350 agtgggatcc caggaattga tgaggtcatg aagactacca aaatcatcat 2400

tgggtgtttt gtggccatca cactcatggc tgcagtgatg ctggtcattt 2450 tctacaagat gaggaagcag caccatcggc aaaaccatca cgccccaaca 2500 aggactgttg aaattattaa tgtggatgat gagattacgg gagacacacc 2550 catggaaagc cacctgccca tgcctgctat cgagcatgag cacctaaatc 2600 actataactc atacaaatct cccttcaacc acacaacaac agttaacaca 2650 ataaattcaa tacacagttc agtgcatgaa ccgttattga tccgaatgaa 2700 ctctaaagac aatgtacaag agactcaaat ctaaaacatt tacagagtta 2750 caaaaaacaa acaatcaaaa aaaaagacag tttattaaaa atgacacaaa 2800 tgactgggct aaatctactg tttcaaaaaa gtgtctttac aaaaaaacaa 2850 aaaagaaaag aaatttatt attaaaaatt ctattgtgat ctaaagcaga 2900 caaaaa 2906

<210> 292

<211> 640

<212> PRT

<213> Homo Sapien

<400> 292

Met Leu Asn Lys Met Thr Leu His Pro Gln Gln Ile Met Ile Gly
1 5 10 15

Pro Arg Phe Asn Arg Ala Leu Phe Asp Pro Leu Leu Val Val Leu 20 25 30

Leu Ala Leu Gln Leu Leu Val Val Ala Gly Leu Val Arg Ala Gln 35 40 45

Thr Cys Pro Ser Val Cys Ser Cys Ser Asn Gln Phe Ser Lys Val
50 55 60

Ile Cys Val Arg Lys Asn Leu Arg Glu Val Pro Asp Gly Ile Ser 65 70 75

Thr Asn Thr Arg Leu Leu Asn Leu His Glu Asn Gln Ile Gln Ile 80 85 90

Ile Lys Val Asn Ser Phe Lys His Leu Arg His Leu Glu Ile Leu
95 100 105

Gln Leu Ser Arg Asn His Ile Arg Thr Ile Glu Ile Gly Ala Phe 110 115 120

Asn Gly Leu Ala Asn Leu Asn Thr Leu Glu Leu Phe Asp Asn Arg 125 130 135

Leu Thr Thr Ile Pro Asn Gly Ala Phe Val Tyr Leu Ser Lys Leu

				140					1 / 5					150
т	C.	т.	TT		70	70	70	_	145	63	G	<b>T</b> 3	_	150
гуѕ	Glu	Leu	Trp	Leu 155	Arg	Asn	Asn	Pro	11e 160	Glu	Ser	Ile	Pro	Ser 165
Tyr	Ala	Phe	Asn	Arg 170	Ile	Pro	Ser	Leu	Arg 175	Arg	Leu	Asp	Leu	Gly 180
Glu	Leu	Lys	Arg	Leu 185	Ser	Tyr	Ile	Ser	Glu 190	Gly	Ala	Phe	Glu	Gly 195
Leu	Ser	Asn	Leu	Arg 200	Tyr	Leu	Asn	Leu	Ala 205	Met	Cys	Asn	Leu	Arg 210
Glu	Ile	Pro	Asn	Leu 215	Thr	Pro	Leu	Ile	Lys 220	Leu	Asp	Glu	Leu	Asp 225
Leu	Ser	Gly	Asn	His 230	Leu	Ser	Ala	Ile	Arg 235	Pro	Gly	Ser	Phe	Gln 240
Gly	Leu	Met	His	Leu 245	Gln	Lys	Leu	Trp	Met 250	Ile	Gln	Ser	Gln	Ile 255
Gln	Val	Ile	Glu	Arg 260	Asn	Ala	Phe	Asp	Asn 265	Leu	Gln	Ser	Leu	Val 270
Glu	Ile	Asn	Leu	Ala 275	His	Asn	Asn	Leu	Thr 280	Leu	Leu	Pro	His	Asp 285
Leu	Phe	Thr	Pro	Leu 290	His	His	Leu	Glu	Arg 295	Ile	His	Leu	His	His 300
Asn	Pro	Trp	Asn	Cys 305	Asn	Cys	Asp	Ile	Leu 310	Trp	Leu	Ser	Trp	Trp 315
Ile	Lys	Asp	Met	Ala 320	Pro	Ser	Asn	Thr	Ala 325	Cys	Cys	Ala	Arg	Cys 330
Asn	Thr	Pro	Pro	Asn 335	Leu	Lys	Gly	Arg	Tyr 340	Ile	Gly	Glu	Leu	Asp 345
Gln	Asn	Tyr	Phe	Thr 350	Cys	Tyr	Ala	Pro	Val 355	Ile	Val	Glu	Pro	Pro 360
Ala	Asp	Leu	Asn	Val 365	Thr	Glu	Gly	Met	Ala 370	Ala	Glu	Leu	Lys	Cys 375
Arg	Ala	Ser	Thr	Ser 380	Leu	Thr	Ser	Val	Ser 385	Trp	Ile	Thr	Pro	Asn 390
Gly	Thr	Val	Met	Thr 395	His	Gly	Ala	Tyr	Lys 400	Val	Arg	Ile	Ala	Val 405
Leu	Ser	Asp	Gly	Thr 410	Leu	Asn	Phe	Thr	Asn 415	Val	Thr	Val	Gln	Asp 420
Thr	Gly	Met	Tyr	Thr	Cys	Met	Val	Ser	Asn	Ser	Val	Gly	Asn	Thr

				425					430					435
Thr	Ala	Ser	Ala	Thr 440	Leu	Asn	Val	Thr	Ala 445	Ala	Thr	Thr	Thr	Pro 450
Phe	Ser	Tyr	Phe	Ser 455	Thr	Val	Thr	Val	Glu 460	Thr	Met	Glu	Pro	Ser 465
Gln	Asp	Glu	Ala	Arg 470	Thr	Thr	Asp	Asn	Asn 475	Val	Gly	Pro	Thr	Pro 480
Val	Val	Asp	Trp	Glu 485	Thr	Thr	Asn	Val	Thr 490	Thr	Ser	Leu	Thr	Pro 495
Gln	Ser	Thr	Arg	Ser 500	Thr	Glu	Lys	Thr	Phe 505	Thr	Ile	Pro	Val	Thr 510
Asp	Ile	Asn	Ser	Gly 515	Ile,	Pro	Gly	Ile	Asp 520	Glu	Val	Met	Lys	Thr 525
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Ala	Val	Met	Leu	Val 545	Ile	Phe	Tyr	Lys	Met 550	Arg	Lys	Gln	His	His 555
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Lys Gly Asn	Gly Thr 275	Ile	Leu	Lys	Gly •	Asp 280	Thr	Gly	Asn	Asn	Asn 285
Trp Ile Pro	Asp Val 290	Gly	Ser	Thr	Trp	Trp 295	Pro	Pro	Lys	Thr	Pro 300
Tyr Ile Pro	Pro Ile 305	Ile	Thr	Asn	Arg	Pro 310	Thr	Ser	Lys	Pro	Thr 315
Thr Arg Pro	Thr Pro	Lys	Pro	Thr	Pro	Ile 325	Pro	Thr	Pro	Pro	Pro 330
Pro Pro Pro	Leu Pro 335	Thr	Glu	Leu	Arg	Thr 340	Pro	Leu	Pro	Pro	Thr 345
Thr Pro Glu	Arg Pro 350	Thr	Thr	Gly	Leu	Thr 355	Thr	Ile	Ala	Pro	Ala 360
Ala Ser Thr	Pro Pro 365	Gly	Gly	Ile	Thr	Val 370	Asp	Asn	Arg	Val	Gln 375
Thr Asp Pro	Gln Lys 380	Pro	Arg	Gly	Asp	Val 385	Phe	Ser	Val	Leu	Val 390
His Ser Cys	Asn Phe 395	Asp	His	Gly	Leu	Cys 400	Gly	Trp	Ile	Arg	Glu 405
Lys Asp Asn	Asp Leu	His	Trp	Glu	Pro	Ile 415	Arg	Asp	Pro	Ala	Gly 420
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Asp Leu Cys	Leu Ser 455	Phe	Arg	His	Lys	Val 460	Thr	Gly	Leu	His	Ser 465
Gly Thr Leu	Gln Val 470	Phe	Val	Arg	Lys	His 475	Gly	Ala	His	Gly	Ala 480
Ala Leu Trp	Gly Arg 485	Asn	Gly	Gly	His	Gly 490	Trp	Arg	Gln	Thr	Gln 495
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Gly Val Arg Gly Gln Ala Leu Tyr Leu Pro Val His Tyr Gly Phe 35 40 45

His Thr Pro Ala Ser Asp Ile Gln Ile Ile Trp Leu Phe Glu Arg
50 55 60

Pro His Thr Met Pro Lys Tyr Leu Leu Gly Ser Val Asn Lys Ser 65 70 75

Val Val Pro Asp Leu Glu Tyr Gln His Lys Phe Thr Met Met Pro  $80 \\ 85 \\ 90$ 

Pro Asn Ala Ser Leu Leu Ile Asn Pro Leu Gln Phe Pro Asp Glu 95 100 105

Gly Asn Tyr Ile Val Lys Val Asn Ile Gl<br/>n Gly Asn Gly Thr Leu 110  $\,$  115  $\,$  120

Ser Ala Ser Gln Lys Ile Gln Val Thr Val Asp Asp Pro Val Thr 125 130 135

Lys Pro Val Val Gln Ile His Pro Pro Ser Gly Ala Val Glu Tyr 140 145 150

Val Gly Asn Met Thr Leu Thr Cys His Val Glu Gly Gly Thr Arg 155 160 165

Leu Ala Tyr Gln Trp Leu Lys Asn Gly Arg Pro Val His Thr Ser 170 175 180

Ser Thr Tyr Ser Phe Ser Pro Gln Asn Asn Thr Leu His Ile Ala 185 190 195

Pro Val Thr Lys Glu Asp Ile Gly Asn Tyr Ser Cys Leu Val Arg 200 205 210

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Phe	Asp	Cys	Ser	Ala 260	Asp	Ser	His	Pro	Pro 265	Asn	Thr	Tyr	Ser	Trp 270
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Ala	Gln	Lys	Gly	Lys 335	Ser	Leu	Ser	Pro	Leu 340	Ala	Ser	Ile	Thr	Gly 345
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Glu	Asp	Ala	Leu	Asp 395	Asp	Phe	Gly	Ile	Tyr 400	Glu	Phe	Val	Ala	Phe 405
Pro	Asp	Val	Ser	Gly 410	Val	Ser	Arg	Ile	Pro 415	Ser	Arg	Ser	Val	Pro 420
Ala	Ser	Asp	Cys	Val 425	Ser	Gly	Gln	Asp	Leu 430	His	Ser	Thr	Val	Tyr 435
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Val Arg Val Ile Thr Asp Glu Asn Trp Arg Glu Leu Leu Glu Gly 35 40 45

Asp Trp Met Ile Glu Phe Tyr Ala Pro Trp Cys Pro Ala Cys Gln 50 55 60

Asn Leu Gln Pro Glu Trp Glu Ser Phe Ala Glu Trp Gly Glu Asp
65 70 75

Leu Glu Val Asn Ile Ala Lys Val Asp Val Thr Glu Gln Pro Gly

Leu Ser Gly Arg Phe Ile Ile Thr Ala Leu Pro Thr Ile Tyr His
95 100 105

Cys Lys Asp Gly Glu Phe Arg Arg Tyr Gln Gly Pro Arg Thr Lys 110 115 120

Lys Asp Phe Ile Asn Phe Ile Ser Asp Lys Glu Trp Lys Ser Ile 125 130 135

Glu Pro Val Ser Ser Trp Phe Gly Pro Gly Ser Val Leu Met Ser 140 145 150

Ser Met Ser Ala Leu Phe Gln Leu Ser Met Trp Ile Arg Thr Cys 155 160 165

His Asn Tyr Phe Ile Glu Asp Leu Gly Leu Pro Val Trp Gly Ser 170 175 180

Tyr Thr Val Phe Ala Leu Ala Thr Leu Phe Ser Gly Leu Leu Leu 185 190 195

Gly Leu Cys Met Ile Phe Val Ala Asp Cys Leu Cys Pro Ser Lys 205 Arg Arg Pro Gln Pro Tyr Pro Tyr Pro Ser Lys Lys Leu Leu 215 220 Ser Glu Ser Ala Gln Pro Leu Lys Lys Val Glu Glu Glu Glu Glu Ala Asp Glu Glu Asp Val Ser Glu Glu Glu Ala Glu Ser Lys Glu 245 250 Gly Thr Asn Lys Asp Phe Pro Gln Asn Ala Ile Arg Gln Arg Ser 260 265 270 Leu Gly Pro Ser Leu Ala Thr Asp Lys Ser <210> 326 <211> 23 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 326 tgaggtgggc aagcggcgaa atg 23 <210> 327 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 327 tatgtggatc aggacgtgcc 20 <210> 328 <211> 21 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 328 tgcagggttc agtctagatt g 21 <210> 329 <211> 25 <212> DNA <213> Artificial Sequence

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<sup>&</sup>lt;211> 533

<212> PRT <213> Homo Sapien

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Val	Ala	Ser	Tyr	Arg 380	Val	Ser	Lys	Ser	Ser 385	Trp	Leu	Glu	Glu	Asp 390
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Thr	Phe	Leu	Asn	Tyr 455	Met	Ser	Asp	Val	Glu 460	Ala	Gly	Gly	Ala	Thr 465
Val	Phe	Pro	Asp	Leu 470	Gly	Ala	Ala	Ile	Trp 475	Pro	Lys	Lys	Gly	Thr 480
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Arg	Thr	Arg	His	Ala 500	Ala	Cys	Pro	Val	Leu 505	Val	Gly	Cys	Lys	Trp 510
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Val Gly Glu Arg Gly Gly Pro Gln Asn Pro Asp Ser Arg Ala Arg
Leu Asp Gln Ser Asp Glu Asp Phe Lys Pro Arg Ile Val Pro Tyr
Tyr Arg Asp Pro Asn Lys Pro Tyr Lys Lys Val Leu Arg Thr Arg
Tyr Ile Gln Thr Glu Leu Gly Ser Arg Glu Arg Leu Leu Val Ala
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Val Leu Thr Ser Arg Ala Thr Leu Ser Thr Leu Ala Val Ala Val
                 110
                                     115
Asn Arg Thr Val Ala His His Phe Pro Arg Leu Leu Tyr Phe Thr
                                     130
Gly Gln Arg Gly Ala Arg Ala Pro Ala Gly Met Gln Val Val Ser
                                     145
His Gly Asp Glu Arg Pro Ala Trp Leu Met Ser Glu Thr Leu Arg
                                     160
His Leu His Thr His Phe Gly Ala Asp Tyr Asp Trp Phe Phe Ile
                 170
                                     175
Met Gln Asp Asp Thr Tyr Val Gln Ala Pro Arg Leu Ala Ala Leu
                 185
                                     190
Ala Gly His Leu Ser Ile Asn Gln Asp Leu Tyr Leu Gly Arg Ala
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                                     205
Glu Glu Phe Ile Gly Ala Gly Glu Gln Ala Arg Tyr Cys His Gly
                 215
Gly Phe Gly Tyr Leu Leu Ser Arg Ser Leu Leu Arg Leu Arg
Pro His Leu Asp Gly Cys Arg Gly Asp Ile Leu Ser Ala Arg Pro
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Ala	Lys	Asn	Arg	Asp 290	Pro	Glu	Lys	Glu	Gly 295	Ser	Ser	Ala	Phe	Leu 300
Ser	Ala	Phe	Ala	Val 305	His	Pro	Val	Ser	Glu 310	Gly	Thr	Leu	Met	Tyr 315
Arg	Leu	His	Lys	Arg 320	Phe	Ser	Ala	Leu	Glu 325	Leu	Glu	Arg	Ala	Tyr 330
Ser	Glu	Ile	Glu	Gln 335	Leu	Gln	Ala	Gln	Ile 340	Arg	Asn	Leu	Thr	Val 345
Leu	Thr	Pro	Glu	Gly 350	Glu	Ala	Gly	Leu	Ser 355	Trp	Pro	Val	Gly	Leu 360
Pro	Ala	Pro	Phe	Thr 365	Pro	His	Ser	Arg	Phe 370	Glu	Val	Leu	Gly	Trp 375
Asp	Tyr	Phe	Thr	Glu 380	Gln	His	Thr	Phe	Ser 385	Cys	Ala	Asp	Gly	Ala 390
Pro	Lys	Cys	Pro	Leu 395	Gln	Gly	Ala	Ser	Arg 400	Ala	Asp	Val	Gly	Asp 405
Ala	Leu	Glu	Thr	Ala 410	Leu	Glu	Gln	Leu	Asn 415	Arg	Arg	Tyr	Gln	Pro 420
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Phe	Asp	Pro	Ala	Arg 440	Gly	Met	Glu	Tyr	Thr 445	Leu	Asp	Leu	Leu	Leu 450
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Pro	Glu	Val	Leu	Asn 605	Arg	Cys	Arg	Met	Asn 610	Ala	Ile	Ser	Gly	Trp 615
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<sup>&</sup>lt;211> 318

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$$35$$
  $40$   $45$ 

Thr Lys His Cys Asp Lys Ala Glu Phe Phe Ser Ser Glu Asn Val
$$95 \hspace{1.5cm} 100 \hspace{1.5cm} 105$$

Cys Leu Lys Tyr Ala Gly Val Phe Ala Glu Asn Ala Glu Asp Ala

a seed to a

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo Sapien

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Pro Asp Phe Phe Leu Leu Leu Phe Arg Gly Cys Leu Ile Gly
20 25 30

Ala Val Asn Leu Lys Ser Ser Asn Arg Thr Pro Val Val Gln Glu Phe Glu Ser Val Glu Leu Ser Cys Ile Ile Thr Asp Ser Gln Thr Ser Asp Pro Arg Ile Glu Trp Lys Lys Ile Gln Asp Glu Gln Thr Thr Tyr Val Phe Phe Asp Asn Lys Ile Gln Gly Asp Leu Ala Gly Arg Ala Glu Ile Leu Gly Lys Thr Ser Leu Lys Ile Trp Asn Val 100 Thr Arg Arg Asp Ser Ala Leu Tyr Arg Cys Glu Val Val Ala Arg Asn Asp Arg Lys Glu Ile Asp Glu Ile Val Ile Glu Leu Thr Val Gln Val Lys Pro Val Thr Pro Val Cys Arg Val Pro Lys Ala Val 140 145 Pro Val Gly Lys Met Ala Thr Leu His Cys Gln Glu Ser Glu Gly His Pro Arg Pro His Tyr Ser Trp Tyr Arg Asn Asp Val Pro Leu Pro Thr Asp Ser Arg Ala Asn Pro Arg Phe Arg Asn Ser Ser Phe 185 190 195 His Leu Asn Ser Glu Thr Gly Thr Leu Val Phe Thr Ala Val His Lys Asp Asp Ser Gly Gln Tyr Tyr Cys Ile Ala Ser Asn Asp Ala Gly Ser Ala Arg Cys Glu Glu Glu Glu Met Glu Val Tyr Asp Leu 235 230 Asn Ile Gly Gly Ile Ile Gly Gly Val Leu Val Val Leu Ala Val Leu Ala Leu Ile Thr Leu Gly Ile Cys Cys Ala Tyr Arg Arg Gly 260 Tyr Phe Ile Asn Asn Lys Gln Asp Gly Glu Ser Tyr Lys Asn Pro 275 280 285 Gly Lys Pro Asp Gly Val Asn Tyr Ile Arg Thr Asp Glu Glu Gly Asp Phe Arg His Lys Ser Ser Phe Val Ile 305 310